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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No.

is the number of results predicted by chance to have a

prophylaxis and detection of cancers

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

SOURCE SOURCE ORGANISM REFERENCE AUTHORS TITLE	AR139128 LOCUS DEFINITION ACCESSION VERSION VERSION	RESULT 1	44 5 5	41	4 3 C	37	, m t	v	31	22 6		c 25	23 24	22	385	18	16 17	15	1 1 1	11	c 10	· 00 ·	<b>J</b> 6	O I		c 2 1	Result No.	
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to 918) iti,P.K. and Kaplan,H.A. iti,P.K. and Kaplan,H.A. ling fragments that specifically encoding the fragments, and use and detection of cancers	918 bp DNA t US 6207153.	ALIGNMENTS	AB064130 HSA245275 HSA245276	HSU80102 HSU43759	HSROU127 AB063793	AB064108 AB021507 WSA226226	AB064107	AB064118	AB064110	AX616568 T08679	AB064067	AR139118 AR139119	AF377321 AX055802						MMVI24FRG		•				AR139130	AR139128 AR139129	ID	Constitution
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Dan,M.D., Maiti,P.K. and Kaplan,H.A.
Antigen binding fragments that specifically
nucleotides encoding the fragments, and use
prophylaxis and detection of cancers
Patent: US 6207153-A 15 27-MAR-2001;
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Pred. No. 2.3e-246;
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1 (bases 1 to 867)

Dan, M.D., Maiti, P.K. and Kaplan, H.A.

Antigen binding fragments that specifically detect cancer cells, and use thereof for the prophylaxis and detection of cancers

Patent: US 6207153-A 16 27-MAR-2001;
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Dan,M.D., Maiti, P.K. and Kaplan,H.A.

Dan,M.D., Maiti, P.K. and Kaplan,H.A.

Antigen binding fragments that specifically detect cancer cells, nucleotides encoding the fragments, and use thereof for the prophylaxis and detection of cancers

Patent: US 6207153-A 18 27-MAR-2001;
                                                                                                                                                                                                                                                                                                          Unclassified
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artificial sequences.

1 (bases 1 to 801)

1 (bases 1 to 801)

Ohba, H., Kashima, K., Tomozawa, T. and Chiba, J

Molecular expression of human antibody and p
antigen-binding structure

(in) PROCEEDINGS OF THE JAPANESE SOCIETY OF
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(Dases 1 to 801)
(Ohba,H., Kashima,K., Tomozawa,T. and Chiba,J.

Direct Submission
Submitted (29-SEP-1997) Department of Biological Sciences and Submitted (29-SEP-1997) Department of Tokyo, 2641 Yamazaki, Noda,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence update by submitter
On Oct 10, 2002 this sequence version
Location/Qualifiers
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Direct Submission
Submitted (10-0CT-2002) Department of Biological Sciences
Technology, Science University of Tokyo, 2641 Yamazaki, No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                278, Japan
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                                                                                              GCCTGGTACCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACC
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AGGGCCACTGGTATCCCAGCCAGGTTCAGTGGCAGTGGGTCTGGGACAGAGTTCACTCTC
                    ACGGCCACTGGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTC
                                                                          GCCTGGTACCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGCTGCATCCACC
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/note="specific to
/codon_start=1
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QPGRSLRLSCVAXSGFTESSYPMTWYCAFFGKGLEWVASISYDGSYKYKVDSMKGRLTI
SRDNSKNTLYLEMNSLTAEDTAVYYCARTAFFNAYDFWGQGTLVTVSS"
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388. .444
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212 c 229 g 179 t
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/protein_id="AAB86468.;
/db_xref="GI:23683340"
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KUFER PETER (DE) ; ZETTL FLORIAN (DE)
BAEUERLE PATRICK A (DE) ; BORSCHERT K
                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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      TEARBLATION="MGMSCIILFLVATATGVHSELQMTQSPSSLSASYGDRVTITCRA
SQSISSYLNMYQQKPGQPPKLLIYMASTRESGVPDRFSGSESGTNYTLTISSLQPEDF
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LYKGGLRGSLYKLKSPLTMMASHYKQHCPPTPETSCATQIITFESFKENLKDFLLVIP
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                                                                                                                                                                                                       /organism="Homo sapiens"
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/db_xref="taxon:9606"
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                                                                                                                                                  /proteIn_id="CAC08835.1"
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                       ORGANISM
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Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; C Mammalia; Eutheria; P

Chordata; Primates;

Craniata; Vertebrata; Eutel Catarrhini; Hominidae; Homo

Euteleostomi;

Sequence 38 AX023367 AX023367.1

GI:10183779 from

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15-SEP-2000

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Query Match
Best Local Similarity
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                  AAAGGGACCACGGTCACCGTCTCCTCAGGATCC 858
                                                                                           GATCAGAGCCTGTTGGGTGACT-----ATGACCACTACGGTTTTGGACGTCTGGGGGC
                                                                                                                                                                 GTGTATCTAAAAATGAACAGCCTGAGAAACTGAGGACACGGCTGTCTATTACTGTGCGAGA
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                                                                        GATATGGGGTGGGCAGTGGCTGGAGACCCTACTACTACTACGGTATGGACGTCTGGGGC
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Patent: WO 006605-A 38 10-FEB-2000;

RUPER PETER (DE); ZETTL FLORIAN (DE); DREIER TORSTEN (DE);

BABUERLE PATRICK A (DE); BORSCHERT KATRIN (DE); MICROMET GES

BIOMEDIZINIS (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kufer, P.,
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                  CAGGCTCTAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCACTAAA
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   CAGGCTCCAGGCAAGGGGCTGGAGTGGCCAGTTATATCATATGATGAAGTAATAAA
                                                              AGACTCTCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGC
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|mol_type="genomic DNA"
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Synthetic construct JB-43
derivative mRNA, partial (
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1 (bases 1 to 867)

Bangale,Y., Li,L., Cavill,D., Gordon,T., Karle,S., Zhou,Y.-X., Taguchi,H., Bhatia,G., Nishiyama,Y., A
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/transl_table=11
/product="recombinant s:
/protein_id="AAL13299.1'
/db_xref="GI:16033615"
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823. .855
                                                         KLISEEDLNGAA"
                                                                            TAVYYCARVGSGWYGGYYYGMDVWGQGTTVTVSSGSASAPTLFPAAAHHHHHHGAAEC
                                                                                                                                                                                                                                                                                                                                                       /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="JB-43"
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                                                                                                                                                                                                                                                                                                                  /cell_type="PBL; B-lymphocyte"
/dev_stage="adult"
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Bangale,Y., Li,L., Cavill,D., Gordon,T., Karle,S.,
Zhou,Y.-X., Taguchi,H., Bhatia,G., Nishiyama,Y., A.
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1 (bases 1 to 861)

Bangale, Y., Li,L., Cavill,D., Gordon,T., Karle,S.,
Zhou,Y.-X., Taguchi,H., Bhatia,G., Nishiyama,Y., Ar
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RLIIYDASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYQQRSNWPLTFGGGT
KLEIKRSSGGGSGGGGGGSGALQVQLQQGGPRLVKPSGTLSLTCTVSGGSISSSSYY
WGWIRQHPGKGLEWICYIYYSGSTYYNPSLKSRVTISVDTSKNQFSLKLSSVTAADTA
VYYCARLQQARGYSYGYGIDYWGQGTLVTVSSGSASAPTLFPAAAHHHHHHGAAEQKL
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/trans[_table=11
/product="recombinant single chain
/protein_id="AAL13298.1"
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817. .849
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/dev_stage="adult"
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                                                                                                                                                      AR139121
Sequence 6 :
AR139121
AR139121.1
      1 (bases 1 to 450)
Dan,M.D., Maiti,P.K. and Kaplan,H.A.
Antigen binding fragments that specifically
nucleotides encoding the fragments, and use
prophylaxis and detection of cancers
Patent: US 6207153-A 6 27-MAR-2001;
Location/Qualifiers
                                                                                                  Unclassified
                                                                                                                 Unknown
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AR139120
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Dan,M.D., Maiti,P.K. and Kaplan,H.A.
Antigen binding fragments that specifically detect cancer cells,
nucleotides encoding the fragments, and use thereof for the
grophylaxis and detection of cancers
Patent: US 6207153-A 4 27-MAR-2001;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (30-MAR-1998) Kalinke U., Institute of Experimental Immunology, Department of Pathology, Schmelzbergetr. 12, 8091 Zuerich, SWITZERLAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      scPv fragment; VI24 antibody.
Mus musculus (house mouse)
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                                                               GATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC 429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GACAGGTCCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTGGAG
                                GACATTGTGATGTCACAGTCTCCATCCTCCCTAGCTGTCAGTTGGAGAGAAGGTTACT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rect Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (bases 1 to 790)
                                                                                                                                                                             183
                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence: X97498.
                                                                                                                                                                         /note="(GlyGlyGlyGlySer)x4 flexible linker"
428. . 778
/note="heavy chain"
191 c 234 g 182 t
                                                                                                                                                                                                                                       /note="light chain" 368. .427
                                                                                                                                                                                                                                                                        /note="DYKD FLAG-tag"
29. .367
                                                                                                                                                                                                                                                                                                                                                /db xref="taxon:10090"
/cell_line="germline precursor
/cell_type="hybridoma"
                                                                                                                                                                                                                                                                                                                                                                                                                   TOW
TOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                    note="VSV-IND neutralising"
                                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                   strain="BALB/c"
                                                                                                            39.0%;
                                                                                                                                                                                                                                                                                                                                                                                                              type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chordata;
Rodentia;
                                                                                             Score 358.4; DB 10;
Pred. No. 1.8e-89;
0; Mismatches 171;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                      8CFV
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                                                                                                                                                                                                                                                                                                                      fragment"
                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                       VI24"
                                                                                                                            Length
                                                                                                                              790;
    -CTACTTAGCC 174
                                                                                               39;
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                                                                                               Gaps
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                                               JOURNAL
MEDLINE
PUBMED
REFERENCE
AUTHORS
                                                                                                                                                                                                                                                                                                                                    RESULT 13
MMVI24FRG
LOCUS
DEFINITION
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SOURCE
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Direct Submission
Submitted (24-APR-1996) U. Kalinke, Institute of Experimental
Immunology, Department of Pathology, Schmelzbergstr. 12, 8091
                                                                                                                                                                                                                                                                                                                                 MMVI24FRG 790 bp DN
M.musculus VI24 antibody VSV-Indiana
                                                                                                                                                          Zinkernagel,R.M. and Hengartner,H.
Monovalent single-chain Fv fragments and bivalent miniantibodies
                                                                                                                                                                                                                                                    scFv fragment; VI24 antibody.
Mus musculus (house mouse)
Mus musculus
                                                                                                             Eur. J. Immunol. 26 (12), 2801-2806 (1996)
                                                                                                                                              bound to vesicular stomatitis
                                                                                                                                                                                            Kalinke, U.,
                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                       X97498.1 GI:1495715
                                                                                                                                                                                                                                                                                                                       X97498
                                                 Kalinke,U
                                                                                                97131797
                                                                                                                              infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAGAGCCTGTTGGGTGACTA 794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TATCTAAAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGAT 774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TACGCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGGTG 714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGGTACCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGG 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGGGTTACTACTTTGACTA 744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TATGCAGACACAGTGAAGGGCCGATTCACCATCTCCAGAGACAATCCCAAGAACACCCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTCAGACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTG 414
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                                                               (bases 1 to
                                                                                                                                                                                      Krebber, A., Krebber, C., Bucher, B.,
                                                               790)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -ATTACACGTTCGGAGGGGGGACCAAGCTGGAAATAAAACGTGGTGGT 373
                                                                                                                                              virus
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                                                                                                                                           protect
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neutralising
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Best Local
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                                                                                                                                                                                                        GCTGCACCATCTGTCTCTGGCGGTGGCGGTTCCCGGAGGTGGATGTGGATCAGGTGGAGGTGGC
                                                                                                                                                                                                                                                                                                                                                GCCACTGGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCCACTCTCACC
                                                                                                                                                                                                                                                                                                                                                                                                       TGGTACCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCATCTATGGTGCATCCACCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                               ATGAGCTGCAAGTCCAGTCAGAGCCTTTTATATAGTAGCAATCAAAAGAACTACTTGGCC 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAG-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGGAAAGAGCCACC 129
              TACGCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGGTG
                                                               GCTCTAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCACTAAATAC
                                                                                                                                                      TCCCAGGTGCAGCTGGTGGAGTCTGGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGA
                                                                                                                                                                                                                                                      CCTCAGACACCTCAGATCACTTTCGGCGGAGGGAGGCCAAGGTGGAGATCAAACGAACTGTG
                                                                                                                                                                                                                                                                                                  ATCAGTAGACTGGAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCA 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GACATTGTGATGTCACAGTCTCCCATCCTAGCTGTCAGTTGGAGAGAAGGTTACT
TATGCAGACACAGTGAAGGGCCGATTCACCATCTCCAGAGACAATCCCAAGAACACCCTG
                                               GCTCCAGAGAGGGGCTGGAGTGGGTCGCATACATTAGTAGTGGCAGTAGTACCATCTTC
                                                                                           CTCTCTGTGCAGCCTCTGGATTCACTTTCAGTAATTTTGGAATGTACTGGGTTCGTCAG
                                                                                                             CTCTCCTGTGCAGCCTCTGGATTCCCCCTTCAGAAGCTTTGCTATGCACTGGGTCCGCCAG
                                                                                                                                                                                         G-----GTGGTTCTGGCGGCGGCGGCTCCGGAGGTGGTGGATCCGGAGGTGGAGGT
                                                                                                                                                                                                                                                                                      ATCAGCAGTGTGAAGGCTGAAGACCTGGCAGTTTATTACTGTCAGCAATATTATAGCT--
                                                                                                                                                                                                                                                                                                                                    GAATCTGGGGTCCCTGATCGCTTCACAGGCAGTGGATCTGGGACAGATTTCACTCTCACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="(GlyGlyGlyGlySer)x4
428. .778
/note="heavy chain"
190 c 233 g 185 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="light chain"
368. .427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="DYKD FLAG-tag"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="spleen"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db xref="taxon:10090"
/cell_line="hybridoma VI24"
/cell_type="B_cell"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic
/strain="BALB/c"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus"
/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note="VSV-IND neutralising scFv fragment"
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71.1%;
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                                                                                                                                                                                                                                       · ATTACACGTTCGGAGGGGGGACCAAGCTGGAAATAAAACGTGGTGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 352; DB 10;
Pred. No. 1.2e-87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches 175;
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Patent: WO 02093519-A 59 21-NOV-2002;
Heintges, Tobias (DE)
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Human high-affinity antibody fragments for essential proteins
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                                                                                                    AGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCTCCTGTG
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                                                                 TAGAGCCTGAAGATTTTGCAGTTTATTACTGTCAGCAGCGG----
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/mol_type="genomic DNA"

/mol_type="genomic DNA"

/db_xref="taxon:32630"

/note="nDNA codierend fur sFv 41 Antikorperfragment"

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0; Mismatches 171;
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Direct Submission
Submitted (25-AUG-1995) David M. Kranz, Biochemistry, U
Illinois, 600 S. Matthews Ave, R.A.L. Bldg, Urbana, IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The antigen-specific, major histocompatibility complex-restricted receptor on T cells. VI. An antibody to a receptor allotype J. Exp. Med. 160 (2), 452-471 (1984) 84291198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct artificial sequences.

1 (bases 1 to 864)
Cho.B.K., Schodin,B.A. and Kranz,D.M.
Characterization of a single-chain antibody to
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864 bp mRNA linear SYN 26-OCT-1995
Synthetic rat anti-T cell receptor b-chain single-chain antibody
mRNA, complete cds.
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Marrack, P.
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3iol. Chem. 270 (43), 25819-25826 (1995)
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monoclonal antibody KJ16-133"
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L 61801, USA
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838 GTCACCGTCTCCTCAGGATCCGAACAAAACTGATCAGCGAAGAAGATCTGAA 890	0	778 AGCCTGTTGGGTGACTATGACCACTACCGTTTGGACGTCTGGGGCAAAGGGACCACG 837	06 CTCCAAATGGACAGTCTGAGATCTGAGGACACGGCCAGTTACTATTGTGCGAGA 7	718 CTAAAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAG 777		58 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGGTGTAT 71	86 CCAAAGAAGGGTCTGGAGTGGGTCGCAACCATCAGTTCTGGTGGTGATGGCACTTACTAT 6	598 CTAGGCAAGGGGCTGGAGTGGCAGTTATATCATATGATGGAAGCACTAAATACTAC 657	26 TCCTGCGCAGCCTCAGGATTCACTTTCAGTAGCTTTCCAATGGCCTGGGTCCGCCAGGCT 5	538 TCCTGTGCAGCCTCTGGATTCCCCTTCAGAAGCTTTGCTATGCACTGGGTCCGCCAGGCT 597	66 GAAGTCCAGCTGCAGGAGTCTGGGGGAGGATTAGTGCAGCCTGGAAGGTCCCTGACAGTC 52	478 CAGGTGCAGCTGGAGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 537	406 AAGAAGGATGCTGCTAAGAAGGATGATGCTAAGAAAGATGCTAAGAAAGA	427 GTCTCTGGCGGTGGCGGTTCCGGAGGTGGATCAGGTGGAGGTGGCTCC 477	346 CCGTACACGTTTGGAGCTGGCACCAAGCTGGAGCTCAAACGGTCCTCCGCGGATGATGCT 405	367 CAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCT 426	298 CAGGTTGAAGATATTGGAACCTATTACTGTCTACAGGTTTCTAGTTCT 345	307 GAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCT 366	238 CCATCAAGGTTCAGTGGCAGTAGATCTGGCACACAGTATTCTCTTAAGATCAACAGACTA 297	247 CCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTG 306	178 AAACCAGGGAAATCTCCTCAGCTCCTGGTCTATAGTGCAACTATCTTGGCAGATGGGGTC 237	187 AAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACTGGCATG 246	121 ACCATCACATGCCAGGCAAGCCAGGACATTGGTACTTCGTTAGTTTGGTATCAGCAG 177	127 ACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAG 186	61 GCCGACGTCCTGGTGACCCAAACTCCTGCCTCCCTGTCTGCATCTCCGGATGAATCTGTC 120	67 GCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCC 126	1 ATGANANAGACAGCTATCGCGATTGCAGTGGCACTGGCCGGCTTCGCTACCGTTGCGCAG 60	7 ATGAAAAAAACCGCTATCGCGATCGCAGTTGCACTGGCTGG	/ Match 37.7%; Score 346.2; DB 12; Length 864; Local Similarity 66.4%; Pred. No. 4.9e-86; Nes 593; Conservative 0; Mismatches 258; Indels 42; Gaps 5;	EQXLISEEDL" 220 a 210 c 220 g 214 t

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AAT08490

Chimeric gene cont plasmid pTERMSC2H1 Single chain antib Human monoclonal a Human monoclonal a

Erythrocyte and Hu Anti-C5 MAb N19/8

n H11 antibody n monoclonal a n monoclonal a

Human H11 antibody C219scFv DNA seque cDNA encoding an a DNA encoding the h

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Coding sequence of
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Single chain bindi
Human interleukin
Human interleukin

Anti-IL8 monoclona Anti-IL8 monoclona

AAT95346
AAT79537
AAC87642
AAV19110
AAV10110
AAV104528
AAF79614
AAAA46898
AAT91614
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Maximum Match 10
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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•	WO9744461-A2.		intron 907 912		/product= nii=80rv construct /product= nii=80rv construct	CDS	Key		Homo sapiens.	Synthetic.					V region; H chain; neoplasia; detection; lymphoma; tumour cell; probe;		Human H11-scFv construct monomer forming DNA sequence.	29-MAY-1998 (first entry)		: AAV10118;		AAV10118 standard; DNA; 918 BP.	AAV10118	RESITAT 1

Result

Score

Query Match

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Description

SUMMARIES

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Best Local Simi
Matches 918;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 918 BP; 213 A; 240 C; 264 G; 201 T; 0 other;
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                                                                         CCATCTGTCTCTGGCGGTGGCGGTTCCCGAAGGTGGTGGATCAAGGTGGAGGTGGCTCCCAAG
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Dan MD,

Maiti PK,

Kaplan HA;

22-MAY-1996;

22-MAY-1997;

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(VIVE-) VIVENTIA BIOTECH INC.

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                                                                                                                                                                                               reading frame; codons"
                                                                                                                                                                                                         /product= "Human H11-single chain variable region (scFv)"
/transl except= (pos:904..915, aa:His-Lys)
/note= "Insertion of two inframe stop codon alters the
reading frame; CDS does not include start and stop
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present DNA sequence encodes human monoclonal antibody (Mab), H11-single chain variable region (scFv). The H11 light chain variable region is linked to the heavy chain variable region through a (SGGGG) linker to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Composition comprising antigen binding fragments of an antibody that recognizes an antigen on neoplastic cells but not on normal cells for
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AAD04538 standard; DNA;

Human Hll-single chain variable region (scFv) complementary DNA

(first entry)

Human; monoclonal antibody; Mab; H11; single chain variable region; scFv; neoplastic disease; melanoma; immunoglobulin IgM; gene therapy; lymphoma; carcinoma; breast; lung; gastric; prostate; ovary; colon; lung; vaccine; neuroblastoma; soft tissue sarcoma; prostatic adenocarcinoma; cytostatic; C-antigen; chronic leukaemia; glioma; ds.

US6207153-B1. sapiens.

27-MAR-2001

240

|CTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTAC

22-MAY-1997; 97US-0862124.

VIVENTIA BIOTECH 푯

WPI; 2001-289584/30. ₹ Maiti PK, Kaplan

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Composition comprising antigen binding fragments of an antibody that recognizes an antigen on neoplastic cells but not on normal cells for use in diagnosis, imaging and treatment of carcinomas

Example 7; Column 65-66; 56pp; English.

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AAD04538/c
ID AAD045
XX AAD045
AC AAD045
XX AAD045
XX Human
XX Human;
XW Human;
XW C-anti
XX Homo 8
XX Hom The present sequence is numan monoconal antibody (Mab) Hil, Hil-(scFv) variable region (scFv) complementary DNA.

The invention relates to human monoconal antibody (Mab) Hil, Hil-(scFv) single chain variable (V) region fragment and their corresponding DNA molecules. Hil antibody is an immunoglobulin of IgM subclass which is specific to C-antigen found specifically on neoplastic cells and not on normal cells. Hil is an antibody obtained from the fusion of peripheral hlood lymphocytes of a 64 year old male with a low grade glioma and fused

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Matches 917
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                                   dimers. This construct is used to determine the ability of H11-ScPv antibody fragments to bind specifically to the C-antigen on cancer cells. Such antigen binding fragments may be used for treating a patient with nepplasia. It is especially useful in the detection of lymphomas and leukaemias where the tumour cells bearing the C antigen are circulating in the patients bloodstream. The polynucleotide sequence may be used as a primer or a probe and the encoded protein may be used in a vaccine or for gene therapy. The human monoclonal antibody (MAb), designated H11, specifically recognises cancerous cells. H11 is specific for glioblastoma, neuroblastoma, malignant melanoma, breast adenocarcinoma, lung adenocarcinoma, small cell lung carcinoma, colon adenocarcinoma and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             H11; monoclonal antibody; MAb; C-antigen; variable region heavy vegion; H chain; neoplasia; detection; lymphoma; tumour cell primer; vaccine; gene therapy; glioblastoma; neuroblastoma; malignant melanoma; adenocarcinoma; small cell lung carcinoma; single chain variable region; scFv; ss.
                                                                                                                                                                                              This sequence encodes a human H11 monoclonal antibody single chain region fragment (H11-scFv) construct which is capable of forming
                                                                                                                                                                                                                                                                   Antigen binding fragment from monoclonal antibody, H11 - tumour specific detection and treatment of neoplasia
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                             adenocarcinoma.
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           CACCATTAGTGAAAGCTT
                                                     ACCGTCTCCTCAGGATCCGAACAAAAACTGATCAGCGAAGAAGATCTGAACCATCACCAT
                                                                                                                                          TGTGCAGCCTCTGGATTCCCCCTTCAGAAGCTTTGCTATGCACTGGGTCCGCCAGGCTCTA
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                                          ACCGTCTCCTCAGGATCCGAACAAAAACTGATCAGCGAAGAAGATCTGAACCATCACCAT
                                                                                    CTGTTGGGTGACTATGACCACTACTACGGTTTGGACGTCTGGGGGCAAAGGGACCACGGTC
                                                                                               CTGTTGGGTGACTATGACCACTACTACGGTTTGGACGTCTGGGGCAAAGGGACCACGGTC
                                                                                                                               AMANTGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGC
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ilarity 94.0%;
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Pred. No. 3.9
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RESULT 5 AAD04539 ID AADO XX

AAD04539

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Query Match Best Local S Matches 863

Similarity

87.1%; nilarity 94.0%; Conservative

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Score 799.6; DB 22; Pred. No. 3.9e-194; 0; Mismatches 4;

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Gaps

Sequence

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other;

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The invention relates to human monoclonal antibody (Mab) H11, H11-(scFv) slingle chain variable (V) region fragment and their corresponding DNA molecules. H11 antibody is an immunoglobulin of IgM subclass which is specific to C-antigen found specifically on neoplastic cells and not on normal cells. H11 is an antibody obtained from the fusion of peripheral blood lymphocytes of a 64 year old male with a low grade glioma and fused to a human myeloma cell line to produce a hybridoma designated NBGM1/H11. A pharmaceutical composition comprising H11 and its derivatives are useful in the diagnosis, imaging and treatment of neoplastic disease, particularly, melanoma, breast carcinoma, lung carcinoma, ovarian carcinoma, colon carcinoma, gastric carcinoma, prostate carcinoma, prostatic adenocarcinoma, B and T cell lymphomas and chronic leukaemias. H11 DNA is also used in vaccines and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neoplastic disease; melanoma; immunoglobulin IdW; gene therapy; lymphoma; carcinoma; breast; lung; gastric; prostate; ovary; colon; lung; vaccine; neuroblastoma; soft tissue sarcoma; prostatic adenocarcinoma; cytostatic; C-antigen; chronic leukaemia; glioma; ds.
                                                                                                                                                                                                                                                                                             The present DNA sequence encodes human monoclonal antibody (Mab), Hll-single chain variable region (scFv). The Hll light chain variable regio is linked to the heavy chain variable region through a (SGGGG)3 linker
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                                                                                                                                                                                                                                                                                                                                                                Example
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                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VIVENTIA BIOTECH INC.
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/transl_except= (pos:853..864, aa:His-Lys)
/note= "Insertion of two inframe stop codon alters the
reading frame; CDS does not include start and stop
codons"
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Human; monóclonal antibody; Mab; H11; single chain variable region; scFv; neoplastic disease; melanoma; immunoglobulin IgM; gene therapy; lymphoma; carcinoma; breast; lung; gastric; prostate; ovary; colon; lung; vaccine; neuroblastoma; soft tissue sarcoma; prostatic adenocarcinoma; cytostatic; C-antigen; chronic leukaemia; glioma; ds.
US6207153-B1
                                                                                                  Homo sapiens.
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22-MAY-1997; 97US-0862124.

22-MAY-1996; 96US-0657449

VIVENTIA BIOTECH INC

PK, Kaplan HA;

WPI; 2001-289584/30

Composition comprising antigen binding fragments of an antibody that recognizes an antigen on neoplastic cells but not on normal cells for use in diagnosis, imaging and treatment of carcinomas -

Column 71-72; 56pp; English

REAR REAR AND SOLUTION OF THE REAR AND SOLUTIO CC variable region (scFv) complementary DNA.

CT whe invention relates to human monoclonal antibody (Mab) H11, H11-(scFv)

CC single chain variable (V) region fragment and their corresponding DNA'

CC supecific chain variable (V) region fragment and their corresponding DNA'

CC specific to C-antigen found specifically on neoplastic cells and not on

CC normal cells. H11 is an antibody obtained from the fusion of peripheral

CC to a human myeloma cell line to produce a hybridoma designated NBGM1/H11.

CC A pharmaceutical composition comprising H11 and its derivatives are

CC useful in the diagnosis, imaging and treatment of neoplastic disease,

CC particularly, melanoma, breast carcinoma, lung carcinoma, ovarian

CC carcinoma, colon carcinoma, gastric carcinoma, prostate carcinoma,

CC lymphoma carcinoma, neuroblastoma, glioma, soft tissue sarcoma, small

CC cell lung carcinoma, prostatic adenocarcinoma, B and T cell lymphomas

CC therary. The present sequence is human monoclonal antibody (Mab), variable region (scFv) complementary DNA. H11-single chain

Sequence 867 BP; 189 A; 241 C; 229 G; 208 T; 0 other;

맑 Ş 문 S 밁 Ş 뮍 Ś 밁 Ś Matches Query Match Best Local : Local Similarity 627 301 241 687 181 747 121 807 867 862; 61 Н AGAGCCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTAC GCGCAGGCCGATATTGTGTTTGACGCAGTCTCCCAGGCACCCCTGTCTTTTGTCTCCAGGGGAA GGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGT 300 AGACTGGAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAG CAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACT GCGCAGGCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAA AGAGCCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTAC Conservative 93.9%; 0, Score 798; DB 22 Pred. No. 1e-193; D; Mismatches 22; ŝ Length Indels 51; Gaps 120 360 180 748 808 568 628 240 60 889

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HD70; single-chain variable fragment; scFv; 17-1A antigen; human; EpCAM; epithelial cell adhesion molecule; inflammatory cytokine; GM-CSF; granulocyte/macrophage colony stimulating factor; heterominibody; CH1-domain; multifunctional compound; heavy chain constant domain; immunoglobulin; cytostatic; immunostimulatory; antileukaemia; diagnosis; antiproliferative; prevention; treatment; malignant; haematopoietic cell; lymphoma; leukaemia; solid tumour; carcinoma; melanoma; sarcoma; ds.
 WO200006605-A2
                                                        misc_feature
                                                                                                                                                                                                                                                                                               HD70scFv-CH1-GM-CSF chain encoding
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/product= "HD70scFv-CH1-GM-CSF
96..842
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                                                                                                   Location/Qualifiers 39..1610
                          /label= HD70_scFv
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produceable in a mammalian host cell as a secretable and fully functional heterodimer of two polypeptide chains, where one of the polypeptide chains comprises, a CHI-domain (constant domain of an immunoglobulin heavy chain) and the other chain comprises CL-domain (constant domain of an immunoglobulin light chain). The polypeptide chains further comprise, fused to the constant domains at least two (poly)peptides having different receptor or ligand functions, where further at least two of the different (poly)peptides lack an intrinsic affinity for one another and are linked via the constant domains. The heterominbodies have cytostatic, immunostimulatory, antileukaemia and antiproliferative activities. These compounds can be used for diagnosing, preventing and treating malignant cell growth related to malignancies of haematopoietic cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas, where the polypeptides are an antiprocess.
                                    The present sequence is a DNA encoding left chain of a heterominibody comprising HD70 single-chain Fv (scFv) fragment N-terminally linked to human CH1 domain which bears at its C-terminus the human inflammatory cytokine granulocyte/macrophage colony stimulating factor (GM-CSF), p. a hexahistidine sequence for ease of purification. HD70 scFv specifically recognises the human epithelial cell adhesion molecule (EpCAM) also called 17-1A antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The patent discloses heterominibodies which are multifunctional compounds produceable in a mammalian host cell as a secretable and fully functional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New multifunctional compounds useful for preventing and/or treating malignant cell growth and for detection and diagnosis -
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                                                                                                     (GM-CSF), plus
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Sequence 1630 BP; 376 A; 484 C; 437 <u>ن</u> 333 T; 0 other;

Matches Query Match Best Local

620;

Conservative

<u>,</u>

Mismatches

145;

Indels

48;

Similarity

48.5%;

Score 445; DB 21; Pred. No. 1.3e-103;

Length 1630;

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                                                                                                                       ACCATCAGTAGACTGGAGCCTGAAGATTTTTGCAGTGTATTACTGTCAGCAGTATGGTAGC
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                                                                TCACCTCAGACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACT
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     WPI; 2000-195265/17
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                                                         Borschert K,
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New multifunctional compounds useful for preventing and/or treating malignant cell growth and for detection and diagnosis \, -
                                                                                                                     AAY44995
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Fig 55B; 166pp; English.

fused to the constant domains at least two (poly) peptides having different receptor or ligand functions, where further at least two of the different (poly) peptides lack an intrinsic affinity for one another and are linked via the constant domains. The heterominibodies have cytostatic, immunostimulatory, antileukaemia and antiproliferative activities. These compounds can be used for diagnosing, preventing and treating malignant cell growth related to malignancies of haematopoietic treating malignant cell growth related to malignancies of haematopoietic The patent discloses heterominibodies which are multifunctional compounds produceable in a mammalian host cell as a secretable and fully functional heterodimer of two polypeptide chains, where one of the polypeptide chains comprises, a CHI-domain (constant domain of an immunoglobulin heavy chain) and the other chain comprises (CI-domain (constant domain of the polypeptide). The present sequence is a DNA encoding right chain of a heterominibody comprising HD70 single-chain Fv (scFv) fragment N-terminally linked to human Ck domain (constant domain of immunoglobulin-kappa light chain) which bears at its C-terminus the human inflammatory cytokine interleukin-2 (II-2). HD70 scFv specifically recognises the human epithelial cell adhesion molecule (EpCAM) also called 17-1A antigen. cells e.g. lymphomas and leukaemias, or to melanomas and sarcomas. an immunoglobulin light chain). The polypeptide chains further solid tumours e.g. carcinomas, two of the comprise,

Sequence 1630 BP; 445 A; 410 C; 408 G; 367 T; 0 other;

Query Match Best Local Similarity Matches 620; Conserv

Conservative

0;

Mismatches 145;

48;

Gaps

48.5%; 76.3%;

Score 445; DB 21,

Length Indels

Ś Ś 밁 S 밁 á 밁 8 밁 Ś 뭐 밁 Ş 밁 뮍 195 315 255 232 112 592 456 472 417 412 52 78 AGGGCCACTGGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTC GCCTGGTACCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACC CCAGGGGAAAGAGCCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTA GCTACCGTTGCGCAGGCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCT 111 ACCATCAGTAGACTGGAGGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGC **AATTGGTATCAGCAGAAACCAGGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACC** GTAGGAGACAGAGTCACCATCACTTGCCGGGCAAGTCAGAG---CATTAGCAGCTATTTA CAGGCTCTAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCACTAAA AGACTCTCCTGTGCAGCCTCTGGATTCCCCTTCAGAAGCTTTGCTATGCACTGGGTCCGC GGCTCCCAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTG GTGGCTGCACCATCTGTCTCTGGCGGTGGCGGTTCCCGGAGGTGGTGGATCAGGTGGAGGT TCACCTCAGACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACT ACCATCAGCAGCCTGCAGCCTGAAGATTTTGCTACTTTTGTCAACAGTCTGACAGT CGGGAATCCGGGTCCCTGACCGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTC GCTACAGGTGTACACTCCGAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCT AGACTOTOCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGC GATCACCTTCGGCCAAGGGACACGACTGGACATTCAA-----GGAGGAGGATCAGGTGGTGGTAGCGGCGGCGGC reerccaeccreeeaecrecere 194 171 137 411 651 575 591 515 531 455 471 416 351 314 291 254 231

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AADO4530/c
ID AADO4530;
XX Human; moth with a single change of the second of the 
cc variable region complementary DNA.

CThe invention relates to human monoclonal antibody (Mab) H11, H11-(scFv) in the invention relates to human monoclonal antibody (Mab) H11, H11-(scFv) in the invention variable (V) region fragment and their corresponding DNA molecules. H11 antibody is an immunoglobulin of IgM subclass which is specific to C-antigen found specifically on neoplastic cells and not on control of the fusion of peripheral control of the fusion of peripheral control of the fusion of peripheral control of the fusion and fused to a human myeloma cell line to produce a hybridoma designated NBGM1/H11.

A pharmaceutical composition comprising H11 and its derivatives are useful in the disgnosis, imaging and treatment of neoplastic disease, particularly, melanoma, breast carcinoma, lung carcinoma, ovarian carcinoma, colon carcinoma, gastric carcinoma, prostate carcinoma, colon carcinoma, gastric carcinoma, prostate carcinoma, small cell lung carcinoma, neuroblastoma, glioma, soft tissue sarcoma, small cell lung carcinoma, prostatic adenocarcinoma, B and T cell lymphomas cand chronic leukaemias. H11 DNA is also used in vaccines and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; monoclonal antibody; Mab; Hl1; light chain variable region; scFv; single chain variable region; immunoglobulin IgM; gene therapy; lymphoma; neoplastic disease; melanoma; carcinoma; breast; lung; gastric; prostate; ovary; colon; lung; chronic leukaemia; neuroblastoma; vaccine; C-antigen; soft tissue sarcoma; cytostatic; prostatic adenocarcinoma; glioma; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Composition comprising antigen binding fragments of an antibody recognizes an antigen on neoplastic cells but not on normal cell use in diagnosis, imaging and treatment of carcinomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Column 59-60; 56pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence is human monoclonal antibody (Mab), H11 light chair
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                                                                                                                                                                                                                                        Human; monoclonal antibody; Mab; H11; light chain variable region; scFv; single chain variable region; immunoglobulin IgM; gene therapy; lymphoma; neoplastic disease; melanoma; carcinoma; breast; lung; gastric; prostate; ovary; colon; lung; chronic leukaemia; neuroblastoma; vaccine; C-antigen; ovary; colon; lung; chronic leukaemia; neuroblastoma; vaccine; C-antigen;
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/transl_except= (pos:259..261, aa:Phe)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           single chain variable (V) region fragment and their corresponding DNA molecules. Hil antibody is an immunoglobulin of IgM subclass which is specific to C-antigen found specifically on neoplastic cells and not on normal cells. Hil is an antibody obtained from the fusion of peripheral blood lymphocytes of a 64 year old male with a low grade glioma and fused to a human myeloma cell line to produce a hybridoma designated NBGM1/Hil. A pharmaceutical composition comprising Hil and its derivatives are useful in the diagnosis, imaging and treatment of neoplastic disease, particularly, melanoma, breast carcinoma, lung carcinoma, olon carcinoma, gastric carcinoma, prostate carcinoma, prostate carcinoma, prostate carcinoma, prostate carcinoma, prostate carcinoma, prostate carcinoma, B and T cell lymphomas and chronic leukaemias. Hil DNA is also used in vaccines and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present DNA sequence encodes human monoclonal antibody light chain variable region.

The invention relates to human monoclonal antibody (Mab) H1 single chain variable (V) region fragment and their corresponding the corresponding to the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Composition comprising antigen binding fragments of an antibody the recognizes an antigen on neoplastic cells but not on normal cells
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Pred. No. 6.2e-82;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1574 BP;
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                                                                                                                                    CDS
                                                                                                                                                                     Mus
                                                                                                                                                                                                   Complement C5; haemolysis; kidney; glomerulonephritis; monoclonal antibody; antiinflammatory; antibody engineering.
                                                                                                                                                                                                                                                             25-MAR-2003
15-MAR-1996
                              02-MAY-1994;
                                                                            09-NOV-1995.
                                                                                                  WO9529697-A1
                                                                                                                                                                                          scFv; single chain antibody;
                                                                                                                                                                                                                                       Anti-C5 MAb
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                                                                                                                                                                                                                                                                                                                     AAT08490
         (ALEX-) ALEXION PHARM INC
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                                                                                                                                                                                                                                      N19/8 scFv (His-tagged)
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(first entry)
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                                                     95WO-US05688
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Matches 552
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Springhorn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MAb, the scFv antibody was unable to block the generation of C5a and C5b-9 from C5. This was in contrast to a humanised scFv (see ARR7616) generated from MAb 5G1.1. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 12; Page 147-149; 181pp; English.
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DB; AAR77617.
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                                                                             TCCAGACACTGTGAAGGGCCGATTCACCATCTCCAGAGACAATGCCAAGAGCACCCTGGA
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TCTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTATTTCTGTGAAGAGA---
                                     TCTAAAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCA 776
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JP, s
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Squinto SP,
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Pred. No. 3e
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Thomas TC, Wang
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Wilkins
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GAGCCTGTTTGGGTGACTATGACCACTACTACGGTTTTGGACGTCTGGGGCCAAAGGGACCAC

836

Query Match Best Local Sim Matches 483;

Similarity

Conservative

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Pred. No. 9.4e-78; 0; Mismatches 164

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Indels Length

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                                                                                                                                                                                                              The present sequence is a chimeric gene containing the cDNA for the immunoglobulin (Ig) heavy and light chain variable regions of a CC murine anti-asparaginase II monoclonal antibody (MAb), and human caparaginase II. The gene was used in the preparation of a novel comprising a biologically active domain and another domain comprising a biologically active domain and another domain containing an epitope, linked via a polypeptide to a 2nd region containing an epitope, linked via a polypeptide to a 2nd region containing an antibody variable region which specifically binds the conformation in which the SCA is bound to the epitope of the 1st region. The chimeric polypeptide assumes a conformation in which the SCA is bound to the epitope of the 1st region and protects its biological activity from deactivation by conformation in which the SCA is bound to the epitope of the 1st region and protects its biological activity from deactivation by conformation in which the scalar activity from deactivation by conformation and protects its biological activity from deactivation by conform agents or alcohol. The regions of the chimeric polypeptide interact to form a structure analogous to an antibody-antigen complex. A L-asparaginase-SCA fusion protein of the above two has better truesin registance than free
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-1988;
31-AUG-1992;
23-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Self-protecting chimeric polypeptide comprising biologically sequence and single-chain antibody sequence - has resistance disrupting temperature, presence of proteolytic enzymes, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric -
Chimeric -
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                                                                                                                                                                                                 antibody-antigen complex. A L-asparaginase-SCA fus
the above type has better trypsin resistance than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ramjeesingh M,
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asparaginase II; monoclonal antibody; MAb; light chain;
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                                                                                                Sequence 1848 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 594;
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                                                                                                                             CACCTCAGACACCTCAGATCACTTTTCGGCGGAGGGGCCCAAGGTGGAGATCAAACGAACTG
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                                         TGGCTGCACCATCTGTCTCTGGCGGTGGCGGTTCCGGAGGTGGATCAGGTGGAGGTG 472
                                                                                             TCCCAC.
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    - GCTGGTGGTGGTTCTGGCGGCGGCGCTCCGGTGGTGGTG

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Pred. No. 1.2e-73;
0; Mismatches 235;
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                                                  1945 AGTCTCCAGAGAAGGGGCTTGAGTGGGTTGCTGAAATTAGATTGAAATCTAATAATTATG 2004
                                                                                                                                                                                                                                                                                                                                                                             1825 GTTCTGAAGTTAAACTGGTCGAGTCTGGAGGAGGATGGTGGAAGCTGGAGGATCCATGA 1884
                                                                                                      2125 GAGGGGTT----------TCATATACTATAGACTACTGGGGTC 2157
2218 TGTA 2221
                                                                                                                                                                                                                                                                                                                                           533 GACTCTCCTGTGCAGCCTCTGGATTCCCCTTCAGAAGCTTTGCTATGCACTGGGTCCGCC 592
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                         887 TGAA 890
                                                                                                                                 767 CGAGAGATCAGAGCCTGTTGGGTGACTATGACCACTACTACGTTTTGGACGTCTGGGGCA 826
                                                                                                                                                                                                                                                                                     593 AGGCTCTAGGCAAGGGGCTGGAGTGGGTGGCAGTTA-----TATCATATGATGGAAGCA 646
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Search completed: September 17, 2003, 03:07:12 Job time : 261 secs

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Regult
No.
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Listing first 45 summaries
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/2
2: /cgn2_6/ptodata/2
3: /cgn2_6/ptodata/2
US-08-657-449-13
918
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Copyright (c) 1993 - 2003 Compugen Ltd
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/BTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfles1.seq:*
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5455030-12

US-08-405-034-3

US-09-240-274-95

US-07-988-430-92

US-08-425-336-89

US-08-488-1138-89

US-08-477-4848-89

US-08-646-360-89

US-08-839-765-89

US-08-839-765-89
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US-08-862-124-15
US-08-862-124-16
US-08-862-124-6
US-08-862-124-6
US-08-862-124-4
US-08-487-2838-19
US-08-447-2828-15
US-08-862-124-1
US-08-862-124-3
US-08-862-124-3
US-08-8611-451-43
US-08-591-632-43
US-08-591-632-51
US-08-765-469-17
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         Sequence 13, Appl
Sequence 16, Appl
Sequence 18, Appl
Sequence 4, Appli
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Sequence 19, Appl
Sequence 11, Appl
Sequence 1, Appli
Sequence 1, Appli
Sequence 3, Appl
Sequence 43, Appl
Sequence 51, Appl
Sequence 51, Appl
Sequence 17, Appl
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Sequence 17, Appl
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Sequence 18, Appl
Sequence 95, Appl
Sequence 95, Appl
Sequence 95, Appl
Sequence 95, Appl
Sequence 89, Appl
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Sequence
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## ALIGNMENTS

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RESULT 1
US-08-862-124-13
                                                                     US-08-862-124-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NOMBER: US/08/862,124
FILING DATE: 22-MAY-1997
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13, Application US/08862124
Patent No. 6207153
GENERAL INFORMATION:
Query Match
Best Local Similarity
                                                                                                                                                                                                                                           TELEX: 706141
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 3160
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                       FEATURE:
NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Dan, Michael D.
APPLICANT: Maiti, Pradip K.
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCOD
TITLE OF INVENTION: FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
TITLE OF INVENTION: DETECTION OF CANCERS
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                      LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                            TELEPHONE: (650) 813-56
TELEFAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Morrison & Foerster LLP STREET: . 755 Page Mill Road CITY: Palo Alto
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join(1..906, 913..918)
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  100.0%;
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  Score 918; DB 3;
Pred. No. 9.8e-252;
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RESULT 2
US-08-862-124-15/c
US-08-862-124-15/c
; Sequence 15, Application U
; Patent No. 6207153
; GENERAL INFORMATION:
; APPLICANT: Dan, Michae
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                                     US/08862124
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US-08-862-124-15
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SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PATENTIN Release #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862
FILING DATE: 22-MAY-1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REGISTRATION NUMBER: 33,943
REGISTRATION NUMBER: 3160
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEPAX: (650) 494-0792
TELEPAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: MOXINISON & F
STREET: 755 Page Mill R
CITY: Palo Alto
STATE: CA
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APPLICANT: Kaplan, F
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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TELEX: 706141
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ZIP: 943
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al Similarity 99.9%;
917; Conservation
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CCATCTGTCTCTGGGGGGGGGGTTCCGGAGGTGGTGGATCAGGTGGAGGTGGCTCCCAG
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22-MAY-1997
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HOWARD A.

HOWARD A.

HOWARD BINDING FRAGMENTS H11, THAT

SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND

DETECTION OF CANCERS
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Pred. No. 2.8e
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RESULT 3
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    NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEFAX: (650) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Maiti, Pradip K.
APPLICANT: Maiti, Pradip K.
APPLICANT: Kaplan, Howard A.
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS Hil, THAT
TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING TITLE OF INVENTION: DETECTION USE THEREOF FOR THE PROPHYLAXIS AND TITLE OF INVENTION: DETECTION OF CANCERS
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS: ADDRESSE: MOTTISON & FOCESTEET LLP
                                                                                                                                                                                      ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                            APPLICATION NUMBER: US/08/862,124
PILING DATE: 22-MAY-1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                    STREET: 755 Page
CITY: Palo Alto
STATE: CA
COUNTRY: USA
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755 Page Mill Road
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iCCATION:
US-08-862-124-16
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Best Local Similarity
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LENGTH: 867 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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Pred. No. 4.7e-218;
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Patent No.
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Matches 862;
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TELEX: 706141
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dan, Michael D.
APPLICANT: Maiti, Pradip K.
APPLICANT: Kaplan, Howard A.
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
TITLE OF INVENTION: DETECTION DETECTION OF CANCERS

NUMBER OF SEQUENCES: 28
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LENGTH: 867 base pairs
TYPE: nucleic acid
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NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862,124
FILING DATE: 22-MAY-1997
CLASSIFICATION: 424
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MEDIOM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Palo Alto
STATE: CA
COUNTRY: USA
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TOPOLOGY: li
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755 Page Mill Road
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Pred. No. 1.4e-217;
0; Mismatches 5;
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                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                 APPLICANT: Dan, Michael D.
APPLICANT: Maiti, Pradip K.
APPLICANT: Maiti, Pradip K.
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING THE
TITLE OF INVENTION: FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
TITLE OF INVENTION: DETECTION OF CANCERS
                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foer
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                        STREET: 755 Page
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94304-1018
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     ARE: PatentIn Release #1.0, APPLICATION DATA:
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US-08-862-124-4
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 450 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.
Matches 361; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEPAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/862,124
FILING DATE: 22-MAY-1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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100.0%; Pred. No. 2.4e-93;
tive 0; Mismatches 0;
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                                             GENERAL INFORMATION:
APPLICANT: Evans, M
APPLICANT: Matis,
APPLICANT: Mueller,
APPLICANT: Nye, Ste
APPLICANT: Rollins,
           APPLICANT:
APPLICANT:
APPLICANT:
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Nye, Steven H.
Rollins, Scott
Rother, Russell P.
Springhorn, Jeremy F
Squinto, Stephen P.

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Evans, Mark J.
Matis, Louis A.
Mueller, Eileen Elliott

US/08487283A

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US-08-862-124-4
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 450 base pairs
TYPE: nucleic acid
STRANDEDINESS: double
TOPOLOGY: linear
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Best Local Similarity
Matches 360; Conserv
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862,124
FILING DATE: 22-MAY-1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
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NAME/KEY:
LOCATION:
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T 433
                                                                       ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC
                                                                                                ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC
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Pred. No. 6.8e-93;
0; Mismatches 1;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/236,
FILLING DATE: 02-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Seth A. Fidel.
REGISTRATION NUMBER: 38,449
REFERENCE/DOCKET NUMBER: ALX-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (203)776-1790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.4M
COMPUTER: Macintosh Cetris:
OPERATING SYSTEM: System 7
SOFTWARE: WordPerfect 3.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (203)772-3655
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Seth A. Fidel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Thomas, Thomas C.
APPLICANT: Wilkins, James R.
TITLE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: OF INFLAMMATORY DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/487,283A FILING DATE: June 7, 1995 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: Nucleic Acid
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CCAGGTGCAGCTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACT 536
                                                                                                                                                                       TGATCCTGTGGAGGCTGATGATGCTGCAACCTATTACTGTCAGCA--
                                                                                                                                                                                                     CAGTAGACTGGAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACC 356
                                                                                                                                                                                                                                        ATCTGGGGTCCCTGCCAGGTTCAGTGGCAGTGGGTCTAGGACAGACTTCACCCTCACCAT
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                                                                TGCACCATCTGTCTCTGGCGGTGGCGGTTCCGGAGGTGGTGGATCAGGTGGAGGTGGCTC 476
                                                                                                       TAATGAGGTTCCGAACACGTTCGGAGGGGGGACCAAGCTGGAAATAAAACGGA-----
                                                                                                                                TCAGACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGC
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N19/8 scFv (His Tag
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                                   CCGGAGGTGGCGGGTCGGGTGGCGGGATCGGGTGGCGGAGGGTC
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Pred. No. 9.5e-89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches 193;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
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US-08-447-422-15
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Patent No. 5686579
                                                                                                                                                           FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/081,410
FILING DATE: 22-UWN-193
APPLICATION NUMBER: US 07/938,505
FILING DATE: 31-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/205,748
FILING DATE: 21-UWN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: SHAMI,
APPLICANT: ROTHSTI
TELEX: 904136
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: RAMJEESINGH, Mohabir
TITLE OF INVENTION: Use of Antibody/Antigen Interactions
TITLE OF INVENTION: Protect or Modulate Biological Activi
                                                       REFERENCE/DOCKET NUMBER: 17
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPHAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                          NAME: BENT, Stephen A. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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RESULT 9
US-08-862-124-1
; Sequence 1, Application US/08862124
; Patent No. 6207153
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; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-447-422-15
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                 GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Mait, Pradip K.
APPLICANT: Mait, Pradip K.
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES E
TITLE OF INVENTION: FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS
TITLE OF INVENTION: DETECTION OF CANCERS
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
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      Morrison & Foerster
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72.3%;
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Pred. No. 3.9e-88;
0; Mismatches 164;
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LS, NUCLEOTIDES ENCODING
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US-08-862-124-3/c
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Sequence 3, Application US/C
Patent No. 6207153
GENERAL INFORMATION:
APPLICANT: Dan, Michael
APPLICANT: Maiti, Pradig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 706141
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 543 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 35.3
Best Local Similarity 89.3
Matches 349; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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LOCATION:
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STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: CA;
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862,124
PILLING DATE: 22-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 424
ATTOMEY/AGENT INFORMATION:
NAME: Lebnhardt, Sussan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 3160
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCTCCTGTGCAGCCTCTGGATTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGGTTTTCCTCGTTGCTCTTTTAAGAGGTATCCAGTGTCAGGTGCAGCTGGTGGAGTCTG
                           ACTACTACGGTTTGGACGTCTGGGGCAAAGG
                                                                                                                      CTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGCCTGTTGGGTGACTATGACC
                                                                                                                                                                                                                    TCACCATCTCCAGAGACACTTCCAAGAACACGGTGTATCTAAAAATGAACAGCCTGAGAA
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ACTACTACGGNTTGGACGCTTGGGGAAAGGG
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Pred. No. 6.7e-83;
0; Mismatches 42;
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Best Local (
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 543 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862,124
FILING DATE: 22-MAY-1997
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kaplan, Howar TITLE OF INVENTION: SPECTIFILE OF INVENTION: FRAGITITLE OF INVENTION: DETE NUMBER OF SEQUENCES: 28 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION UNMER: 33,943
REFERENCE/DOCKET NUMBER: 3160
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEFAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 755 Page Mill Road CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
les 349; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: double
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                                 ACTACTACGGTTTGGACGTCTGGGGCAAAGG 830
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                                                                                                    CTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGCCTGTTGGGTGACTATGACC 799
                                                                                                                                                             TCACCATCTCCAGAGACACTTCCAAGAACACGGTGTATCTAAAAATGAACAGCCTGAGAA 739
                                                                                                                                                                                                             TGGCAGTTATATCATATGATGGAAGCACTAAATACTACGCAGACTCCGTGAAGGGGCGAT 153
                                                                                                                                                                                                                                                                                   CCTTCAGAAGCTTTGCTATGCACTGGGGTCCGCCAGGCTCTAGGCAAGGGGGCTGGAGTGGG
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                                                                   {	t CTGAGGACACGGCTGTCTTTTACTTGTGCGAAAGACAGAGCCTGCTGGGGTGACTATGACCC}
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WENTION: ANTIGEN BINDING FRAGMENTS H11, THAT

VENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING

VENTION: FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND

VENTION: DETECTION OF CANCERS

EQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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Pred. No. 6.7e-83;
0; Mismatches 42
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RESULT 11 US-08-591-632-43

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                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/308,841

FILING DATE: 19-SEP-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/233,619

FILING DATE: 26-APR-1994

PRIOR APPLICATION NUMBER: US 08/139,409

FILING DATE: 19-OCT-199

APPLICATION NUMBER: US 08/139,409

FILING DATE: 19-OCT-199

ATTORNEY/AGENT INFORMATION:

NAME: Fitting, Thomas

REGISTRATION NUMBER: 34,163

REFERENCE/DOCKET NUMBER: TSRI 332.3

TELEPHONE: (619) 784-2937
                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (619) 784-9399
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 4691 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patenth Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lerner, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: sing
TOPOLOGY: circular
MOLECULE TYPE: DNA (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/591,632
FILING DATE: 19-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/11907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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                               ANACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACTGGCATG 246
                                                                                                                                     GCGG----CCGAGCTCACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCC
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                                                                                                                                                                                                      ACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAG
                                                                                                                                                                                                                                      Application US/08591632
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10550 No. 6261558th Torrey Pines Road, TPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                            DNA (genomic)
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                                                                                                                                                                                                                                                                                      34.7%;
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Righard A.
SYNTHETIC
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Pred. No. 8.7e-81;
0; Mismatches 31;
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                                                                                                                                                                                                                                                                                                        Length 4691;
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RESULT 12
US-09-611-451-43
J Sequence 43, Application US/09611451
Patent No. 6395275
GENERAL INFORMATION:
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MOLECULE TYPE: DNA (genomic)
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SEQUENCE DESCRIPTION: SEQ ID
US-09-611-451-43
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Query Match
Best Local Similarity
Matches 375; Conserv
                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 43:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/611,451
FILING DATE: 06-Jul-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/591,632
FILING DATE: 2001-10-29
APPLICATION NUMBER: US 08/308,841
FILING DATE: 19-SEP-1994
APPLICATION NUMBER: US 08/233,619
FILING DATE: 26-APR-1994
APPLICATION NUMBER: US 08/233,619
FILING DATE: 19-OCT-193
ATTORNEY, AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                       NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: TSRI 332.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 784-2937
TELEPAX: (619) 784-9399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Barbas, Carlos F.
Burton, Dennis R.
Lerner, Righard A.
TITLE OF INVENTION: SYNTHETIC HUMAN NEUTRALIZING MONOCLONAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3013
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                                                                                                                                            LENGTH: 4691 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent Counsel

STREET: 10550 No. 6395275th Torrey Pines Road, TPC
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: CA
COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: The Scripps Research Institute, Office of
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                 34.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTIBODIES TO HUMAN IMMUNODEFICIENCY VIRUS
Score 318.4; DB 4
Pred. No. 8.7e-81;
0; Mismatches 31
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                                    DB 4;
                                   Length 4691;
   Indels
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US-08-591-632-51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 51, Applicat Patent No. 6261558 GENERAL INFORMATION:
                           COMPUTER: IBM PC COMPACTION
COMPUTER: IBM PC COMPACTION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/591,632
FILING DATE: 19-CCT-1994
PRIOR APPLICATION NUMBER: PCT/US94/11907
FILING DATE: 19-CCT-1994
PRIOR APPLICATION UNMBER: US 08/308,841
APPLICATION UNMBER: US 08/308,841
FILING DATE: 19-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/33,619
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/233,619
FILING DATE: 26-APR-1994
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/233,619
PRIOR APPLICATION DATA:
IN OR APPLICATION DATA:
APPLICATION UNMBER: US 08/233,619
PRIOR APPLICATION DATA:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                             ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1V-1
CITY: La Jolla
TMATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Borton, Dennis R.
APPLICANT: Lerner, Righard A.
TITLE OF INVENTION: SYNTHETIC:
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: The SCILEFY
ADDRESSEE: Patent Counsel
ADDRESSEE: 10550 No. 6261558th Torrey Pines Road, TPC
 APPLICATION NUMBER: US 0 FILING DATE: 19-OCT-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The Scripps Research Institute, Office of
                                                                                                                                                                                                                                                                                                                                       Floppy disk
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SYNTHETIC HUMAN NEUTRALIZING MONOCLONAL
SYNTHETIC HUMAN IMMUNODEFICIENCY VIRUS
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Dennis R.
                 US 08/139,409
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126

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RESULT 14
US-09-611-451-51
IS-09-611-451-51
Sequence 51, Application US/09611451
Patent No. 6395275
GENERAL INFORMATION:
APPLICANT: Barbas, Carlos F.
Burton, Dennis R.
Burton, Dennis R.
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INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 6166 base pairs
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REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: TSRI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 784-2937
TELEPAX: (619) 784-9399
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: circular
                                                                                                                                                                                                    TITLE OF INVENTION: SYNTHETIC HUMAN NEUTRALIZING MONOCLONAL ANTIBODIES TO HUMAN IMMUNODEFICIENCY VIRUS
      COMPUTER READABLE FORM:
                                                                                                                                                                    CORRESPONDENCE
                                                                                                                                                                                     NUMBER OF SEQUENCES: 92
                                        CITY: La Joll
STATE: CA
COUNTRY: USA
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ADDRESSEE: The Scripps Research Institute,
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                                                                                  Jolla
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                                                                                                                        Patent
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                                                                                                                                             Office of
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                  RESULT 15
US-08-442-542-17
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Sequence 17, Application US/08442542
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/591,632
FILING DATE: 2001-10-29
APPLICATION NUMBER: US 08/308,841
FILING DATE: 19-SEP-1994
APPLICATION NUMBER: US 08/233,619
FILING DATE: 26-APR-1994
APPLICATION NUMBER: US 08/139,409
FILING DATE: 19-0CT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 6166 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: circular
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID N
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                                                                                                     4943
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REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: TSRI 332.3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 784-2937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER; US/09/611,451
FILING DATE: 06-Jul-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4658 ACCCTCTGCAGGGCCAGTCACAGTGTTAGCAGGGCCTACTTAGCCTGGTACCÁGCAG
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                                                                                                                                           427
                                                                                                                                                                                                                       367
                                                                                                                                                                                                                                                                                                                                                                                     247
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                                                                                                                                                                                                                                                                                                       GAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCT 366
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
                                                                                                                                           430
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      34.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NO: 51:
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Gaps

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Query Match Best Local S Matches 375

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; FEATURE:
; NAME/KEY: CDS
; LOCATION: 1..1797
; OTHER INFORMATION:
; OTHER INFORMATION:
US-08-442-542-17
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Best Local Similarity
Matches 493; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 1797 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PILING DATE: 28-JUN-1994
ATTORNEY AGENTI INFORMATION:
NAME: SPELLII, W. MULTRAY
REGISTRATION NUMBER: 32,943
REFRENCE/DOCKET NUMBER: CGC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/442,542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Carozzi, Nadine B.
APPLICANT: Koziel, Michael G.
TITLE OF INVENTION: Antibodies which Bind to
TITLE OF INVENTION: Proteins and their Use
NUMBER OF SEQUENCES: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 16-MAY-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Ciba-Geigy Corporation
STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Hawthorne
STATE: NY
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
                                                    307
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                                                                                                                                                                                                       181 CAGCAGAAACCTGGCCCAGGCTCCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACT
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                    361
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ACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCA
                                                                                                                                         GGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGT
                                                    CCTATGGAGGAGGATGATACTGCAATATATTTCTGTCAGCA------AAGTAGG
                                                                                 AGACTGGAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAG
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from pCIB4631"
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                                                                                                                                                                                                                                                                                                                                                                                     Score 303.6; DB 1;
Pred. No. 9.3e-77;
0; Mismatches 189;
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                                                                       TATEGCEATTCAGTGAAAGACAGGTTCACCGTCTCCAGAGATGATTCACAAAGCATGTTC
                                                                                                                                              GGAAAGGGTTTGGAATGGGTTGCTCGCATAAGAAGTAAAAGTAATAATTATGCAACATCT
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                     TATCTAAAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAG
                                                                                                   TACGCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGGTG
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Search completed: September 17, 2003, 04:23:14
Job time : 68 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Published Applications NA:*

1: /cgn2_6/ptodata/1/pubpna/US07 PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US06 NEW PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06 NEW PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US07 NEW PUB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US07 NEW PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08 NEW PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08 NEW PUB.seq:*

7: /cgn2_6/ptodata/1/pubpna/US08 PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US08 PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US09E PUBCOMB.seq:*

10: /cgn2_6/ptodata/1/pubpna/US09E PUBCOMB.seq:*

11: /cgn2_6/ptodata/1/pubpna/US09E PUBCOMB.seq:*

12: /cgn2_6/ptodata/1/pubpna/US10B PUBCOMB.seq:*

13: /cgn2_6/ptodata/1/pubpna/US10B PUBCOMB.seq:*

14: /cgn2_6/ptodata/1/pubpna/US10B PUBCOMB.seq:*

15: /cgn2_6/ptodata/1/pubpna/US10B PUBCOMB.seq:*

16: /cgn2_6/ptodata/1/pubpna/US10B PUBCOMB.seq:*

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17: /cgn2_6/ptodata/1/pubpna/US10B.seq:*
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 US-09-782-397-13

US-09-782-397-15

US-09-782-397-18

US-09-782-397-18

US-09-782-397-1

US-09-782-397-1

US-09-782-397-3

US-09-782-397-3

US-09-782-397-3

US-09-782-397-3

US-09-782-397-3

US-09-782-397-3

US-09-782-397-3

US-09-782-397-3

US-09-783-324-49

US-09-783-234-37

US-09-783-234-41

US-09-783-234-71

US-09-783-234-71

US-09-783-234-71

US-10-325-694-143
                                                                                                                                                                                                                                                                                                   Description
              Sequence 15,
Sequence 16,
Sequence 6, A
Sequence 1, A
Sequence 3, A
Sequence 16,
Sequence 16,
Sequence 27,
Sequence 37,
Sequence 37,
Sequence 71,
Sequence 71,
Sequence 71,
Sequence 72,
Sequence
                                                                                                                                                                                                                                                                  Sequence 13,
 13, Appl
16, Appl
18, Appl
18, Appl
18, Appl
4, Appl
1, Appl
49, Appl
41, Appl
41, Appl
41, Appl
41, Appl
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US-09-782-397-13

Sequence 13, Application US/09782397

Publication No. US20030021779A1

GENERAL INFORMATION:

APPLICANT: Dan, Michael D.

APPLICANT: Maiti, Pradip K.

"---ard A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  വ
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APPLICATION UNMBER: 08/862,124

FILING DATE: 1997-05-22

ATTORNEY/AGENT INFORMATION:

NAME: Lehnhardt, Susan K.

REGISTRATION NUMBER: 33,943

REFERENCE/DOCKET NUMBER: 31608-20001.20

TELECOMMUNICATION INFORMATION:

TELEPHONE: (650) 813-5600
                                                                                                                                                                       ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION UNMERR: US/09/782,397
FILING DATE: 13-Feb-2001
CLASSIFICATION: CUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                             STREET: 755 Page Mill Road CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                         STATE: CA
                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Maiti, Pradip K.

Kaplan, Howard A.

OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT

SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING

FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
     (650)
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US-10-041-860-67
US-09-453-234-79
US-09-453-234-51
US-10-256-615-3
US-09-453-234-75
US-09-453-234-75
US-09-453-234-77
US-09-453-234-77
US-09-453-234-77
US-10-256-615-3
US-10-256-615-3
US-10-198-846-13125
US-10-330-613-31
US-10-330-613-31
US-10-386-613-31
US-10-386-646-66
US-10-158-646-66
US-10-158-646-67
US-10-158-646-69
US-10-158-646-798-95
US-10-158-646-57
US-09-948-798-194
US-10-158-646-57
US-09-948-798-194
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Sequence 67, Appl
Sequence 33, Appl
Sequence 31, Appl
Sequence 51, Appl
Sequence 62, Appl
Sequence 37, Appl
Sequence 43, Appl
Sequence 41, Appl
Sequence 31, Appl
Sequence 31, Appl
Sequence 13125, A
Sequence 60, Appl
Sequence 64, Appl
Sequence 66, Appl
Sequence 67, Appl
Sequence 99, Appl
Sequence 99, Appl
Sequence 99, Appl
Sequence 97, Appl
Sequence 97, Appl
Sequence 98, Appl
Sequence 57, Appl
Sequence 57, Appl
Sequence 57, Appl
Sequence 230, App
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Match

Length DB

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100.0 99.8 86.9 39.2 39.2 39.3 34.8 33.8 33.8 33.8 33.8 33.8

918 918 867 867 450 450 450 678 678 678 678 678

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seq length: 0 seq length: 2000000000

Title: Perfect score:

US-08-657-449-13 918

OM nucleic

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Scoring table: Sequence:

IDENTITY\_NUC Gapop 10.0 , Gapext 1.0

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LUCATION: join(1...906, 913...918);
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-782-397-13
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INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
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Local Similarity 100.0%;
hes 918; Conservation-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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LOCATION:
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                CTGTTGGGTGACTATGACCACTACTACGGTTTGGACGTCTGGGGCAAAGGGACCACGGTC
                                                                      AAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGC
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Pred. No. 7.6e-270;
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RESULT 2
US-09-782-397-15/c
US-09-782-397-15/c
; Sequence 15, Application US/09782397
; Publication No. US20030021779A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
Michael D.
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Best Local Similarity
Matches 917; Conser
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/782,397

FILING DATE: 13-Feb-2001

CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,124

FILING DATE: 1997-05-22

ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSE: Morrison & Foerster LLP
STREET: 755 Page Mill Road
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                        GCGCAGGCCGATATIGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCCAGGGGAA 120
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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COUNTRY: USA
                    AGAGCCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTAC
                                                                                                                                        LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
 AGAGGCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTAC
                                                                    GCGCAGGCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                            99.8%;
nilarity 99.9%;
Conservative
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Kaplan, Howard A.
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Pred. No. 2.4e-269;
0; Mismatches 1;
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RESULT 3
US-09-782-397-16
; Sequence 16, Application US/09782397
; Publication No. US20030021779A1
; GENERAL INFORMATION:
; APPLICANT: Dan, Michael D.
; Maiti, Pradip K.
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                                                                            NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
                                                                                                                                                            TITLE
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ADDRESSEE: Morrison & F
STREET: 755 page Mill H
CITY: Palo Alto
STATE: CA
COUNTRY: USA
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                                                                                                         Maiti, Pradip K.
KAPJAN, HOWARD A.
KAPJAN, HOWARD FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                Road
                                                               Foerster LLP
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/782,397
FILING DATE: 13-Feb-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY/ACENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PETENTIN Release #1.0,
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TYPE: nucleic acid
STRANDEDNESS: single
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nilarity 94.0%;
Conservative
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Pred. No. 1e-233;
0; Mismatches 4;
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NO: 16:
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                           TELEX: 706141
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                             APPLICATION NUMBER: US/09/782,397
FILING DATE: 13-Feb-2001
CLASSIFICATION: -CUNKnown>
PRIOR APPLICATION DATA: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                    ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Dan, Michael D.
Malti, Pradip K.
Malti, Pradip K.
Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DEFFECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                      NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison &
STREET: 755 Page Mill
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STATE: CA
COUNTRY: USA
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LENGTH: 867 base pairs
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                                                                           TELEFAX: (650) 494-0792
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Pred. No. 3.2e-233;
0; Mismatches 5;
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No
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                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 361; Conserv
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Publication No. US20030021779A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 706141
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPAtible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/782,397
FILING DATE: 13-Feb-2001
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster LLP
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maiti, Pradip K.
KADlan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING THE
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
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                                      310
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                                                                                                 250 GACAGGTTCAGTGGCAGTGGGTCCGGGGACAGACTTCACTCTCACCCATCAGTAGACTGGAG
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                                                                                                                                                                                                                                                            130 CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAGAAA 189
                                                                                                                                                                                                                                                                                                               378 GATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC
                                                                                                                                                                                                                                                                                                                                                      70 GATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC
                        CCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCTCAG
                                                                                                                                                                                                                                      CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAGAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 450 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                           GACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTGGAG
                                                                                                                                                        CCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACTGGCATGCCA
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CCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCTCAG
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                                                                                                                                                                                                                                                                                                                                                                                          39.3%; Score 361; ilarity 100.0%; Pred. No. 6 Conservative 0; Mismatche
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6e-100;
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RESULT 6
US-09-782-397-4
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                                                                                                                                                                   Matches 360;
                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 706141
INFORMATION FOR SEQ ID NO: 4:
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                                                                                                                                                                                                                                                                                                      FEATURE: CDS
                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 450 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/782,397
FILING DATE: 13-Peb-2001
CLASSIFICATION: CURKNOWN>
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Maiti, Pradip K.

Kaplan, Howard A.

TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT

SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING THE
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                                                                                   LOCATION: 1..450
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS
CORRESPONDENCE MORTISON & FOORSTOR LLP
STREET: 755 Page Mill Road
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                                                                                73 GATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC
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                          CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAGAAA
CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAGAAA
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                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Palo Alto
                                                                                                                                                                   Conservative
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                                                                                                                                                                                    Score 359.4; DB 1
Pred. No. 1.9e-99;
                                                                                                                                                                   Mismatches
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                                                                                                                                                                   Indels
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 US-09-782-397-1
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US-09-782-397-1
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Publication No. US20030021779A1
GENERAL INFORMATION:
APPLICANT: Dan, Michael D.
                                                                                                                                                          TELEX: 706141
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kaplan, Howard A.

TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT

SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                  SEQUENCE DESCRIPTION:
                                                                      FEATURE:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison
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                                   NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/782,397
FILING DATE: 13-Feb-2001
CLASSIFICATION: CUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: CA
COUNTRY: USA
ZIP: 94304-1018
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                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC
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                                                                                       TOPOLOGY:
                                                                                                                                       LENGTH: 543 base pairs
                                                                                                                                                                                                              TELEPHONE: (650) 813-56
TELEFAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Maiti, Pradip K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25: Morrison & 755 Page Mill
                                                                                   linear
                                                    CDS
                  SEQ ID
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Road
                  NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-782-397-3/c; Sequence 3, Application US/09782397; Publication No. US20030021779A1; GENERAL INFORMATION:
APPLICANT: Dan, Michael D.; APPLICANT: Dan, Michael D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 8
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           NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,124
APILING DATE: 1997-05-22
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSE: MORTISON & FOETSTET LLP
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Maiti, Fradip K.
Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES E
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS
DETECTION OF CANCERS
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                                                                                                                                                                         APPLICATION NUMBER: US/09/782,397
FILING DATE: 13-Feb-2001
CLASSIFICATION: <Unknown>
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(650)
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Sequence 16, Application US/10153382
Publication No. US20030086930A1
GENERAL INFORMATION:
APPLICANT: PFIZER PRODUCTS INC.
TITLE OF INVENTION: USES OF ANTI-CTLA-4 ANTIBODIES
FILE REFERENCE: PC23019A
CURRENT APPLICATION NUMBER: US/10/153,382
CURRENT FILING DATE: 2002-05-22
PRIOR APPLICATION NUMBER: 60/293042
PRIOR APPLICATION NUMBER: 60/293042
PRIOR APPLICATION NUMBER: 60/293042
PRIOR PILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 16
LENGTH: 1413
TYPE: DNA
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US-10-153-382-16
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Best Local
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ORGANISM: Homo
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
500 GGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCTCCTGTGCAGCCTCTGGATTCC
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TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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Pred. No. 3.3e-87;
0; Mismatches 62
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Pred. No. 9.5e-89;
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US-09-453-234-49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 49, Application US/09453234
Publication No. US20030091995A1
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 49
                                                                                                                                                                                                                                                                                                                                                                                            Matches 338;
                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Lonberg, Nils
APPLICANT: Biogite Diagnostics, Inc.
APPLICANT: Genpharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Ni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA .
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: (1)..(678)
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                                                                                 GACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTGGAG 309
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                                                                                                                                                                                                                                                                                                      GAAATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC
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93.6%;
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Pred. No. 1.7e-84;
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; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(678)
; OTHER INFORMATION: M2-33L
US-09-453-234-85
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US-09-453-234-85
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Best Local Similarity
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: GenPharm International TITLE OF INVENTION: Human Antibodies FILE REFERENCE: 020015-000110US CURRENT APPLICATION NUMBER: US/09/453,234 CURRENT FILING DATE: 1999-12-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
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TYPE: DNA
ORGANISM: Homo sapiens
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           ; Sequence 41, Application US/09453234
; Publication No. US20030091995A1
; GENERAL INFORMATION:
   APPLICANT: Buechler, Joe
   APPLICANT: Valkirs, Gunars
; APPLICANT: Gray, Jeff;
; APPLICANT: Lonberg, Nils
                                                                                                                                      RESULT 13
US-09-453-234-41
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APPLICANT:
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Gray, Jeff Lonberg, Nils Biosite Diagnostics, Inc.

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US-09-453-234-37
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US-09-453-234-37
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnosti
APPLICANT: GenPharm Internat
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Publication No.
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Best Local
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FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
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352
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                                                                                                          370 ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC
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Biosite Diagnostics, I
GenPharm International
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Pred. No. 4.9e-83;
0; Mismatches 17;
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129

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APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Gray, Jeff
APPLICANT: Gray, Jeff
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
ITITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTMARE: PatentIn Ver. 2.1
SEQ ID NO 71
LENCTH: 678
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ITITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 41
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                                                                                                                                                                                                                                                                                                                Sequence 71, Application US/09453234 Publication No. US20030091995A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 335; Conserv
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(678)
OTHER INFORMATION: M1-5L
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Pred. No. 4.9e-83;
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GENERAL INFORMATION:

APPLICANT: Corvalan, Jose R.F.

APPLICANT: Jia, Xiao-Chi

APPLICANT: Peng, Xiao

APPLICANT: Yang, Xiao-Dong

APPLICANT: Yang, Xiao-Dong

APPLICANT: Chen, Francine

APPLICANT: Weber, Richard

APPLICANT: Weber, Richard

APPLICANT: Bezabeh, Binyam

TITLE OF INVENTION: ANTIBODIES DIRECTED TO PDGFD AND USES

TITLE OF INVENTION NUMBER: US/10/041,860
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; NAME/KEY: CDS
; LOCATION: (1)..(678)
; OTHER INFORMATION: M2-11L
US-09-453-234-71
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                                                                                                                                                                SEQ ID NO 59
LENGTH: 379
                                                                Query Match
Best Local
                                                    Matches 339;
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Best Local Similarity
Matches 335; Conserv
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NUMBER OF SEQ ID NOS: 377
                                                                                                                                                                                                SOFTWARE: FastSEQ
                                                                                                                              TYPE: DNA
ORGANISM: homo sapiens
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Pred. No. 1.6e-82;
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Search completed: September 17, 2003, 05:31:54 Job time : 876 secs

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AW401728 UI-HF-BK0
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68552	552	10	707	.2	299.8	43
957976	5797	14	718		8	42
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7917	579	13	1201	'n	302.4	
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BQ707037 AGENCOURT	BQ707037	13	974	ω	303	
11051	BQ711051	13	N	ω	0	
01189	BX368110	13	ū	·	0	
H-IU	AW405817	ø	4	ω.	0	
5216 I	AW405216	9	4	ω	0	
F91729	BF917294	10	0	ω	0	
M830796	BM830796	12	8	ω	0	
v	BF914740	10	4	ω	0	
F914735	BF914735	10	441	ω ·	30	
F663521	BF663521	10	1153	ω ·	3.	
**	BM919414	12	1069	ω ·	<u>မ</u>	
Q331002	BQ331002	13	526	·	04.	
1272 A	BQ711272	13	933	ω.	304.6	
910 1	AW404910	φ.	603	ω	05.	
AW405697 UI-HF-BL0	AW405697	ဖ	472	ω ·	05.	21
451	AW406451	9	456	·	30	
8339	183	12	1014	ω.		
8749	75874	10	862	ω ·	•	18
576	76	9	518	ω.	306.2	17
9298	992	13	801	·	.7	16
09245	0924	13	998	u !	7	15
6634	7066	<u>۔</u>	926	س	.7	14
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686748	5867	5	734	u	.7	12
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74079 IL3-ET01	407	10	N.	33.7	30	σ,
Q709399	Q7093	13	7			U.

a

## BF664453 602146204F1 NIH\_MGC\_48 Homo sapiens cDNA clone IMAGE:4309680 Homo sapiens BF664453.1 GI:11938258 EST. BF664453 mRNA sequence. Homo sapiens (human)

SOURCE ORGANISM VERSION KEYWORDS RESULT 1
BF664453
LOCUS
DEFINITION

EST 21-DEC-2000 3:4309680 5',

ALIGNMENTS

ACCESSION

REFERENCE AUTHORS Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.B. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.B. Consortium/LLNL at:

http://image.llnl.gov

Plate: LLCM1183 row: h column: 01 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 584)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Plate: LLCM1183 row: h column: 01 High quality sequence stop: 584. Unpublished

COMMENT

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TITLE
JOURNAL
COMMENT
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AW401728
LOCUS
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AUTHORS
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Unpublished
Contact: Robert Strausberg,
Email: cgapbs-r@mail.nih.go
Eco RI site shown at the be
                                                       Eukaryota; Metazoa; Chordata; Craniata; V
Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 516)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                               516 bp
UI-HF-BKO-aaf-f-12-0-UI.r1 NIH N
IMAGE:3053711 5', mRNA BECTIONICA
AWA01770
                                                                                                                                     Homo
                                                                                                                                                                           AW401728
AW401728.1 GI:6920414
                                                                                                                                                Homo sapiens (human)
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                                                                                                                                   sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="primary B-cells from tonsils (cell line)"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/note="Porgan: B-cells; Vector: pOTB7; Site_1: XhoI;
/site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."

39 a 168 c 153 g 124 t
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 2.7e-71;
0; Mismatches 44
     beginning
                                   Ph.D.
                                                                                                       Craniata; Vertebrata; Catarrhini; Hominidae;
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                                              AW402624 536 bp mRNA
UI-HF-BKO-aax-h-08-0-UI.rl NIH MGC_36 HC
IMAGE:3055359 5', mRNA sequence.
AW402624
                                AW402624.1
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                                                                                                                                                             CC 514
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     sapiens (human)
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/db_xref="taxon:9606"
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                                 GI:6921329
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Tissue Procurement: Louis M. Staudt, M.D., P
CDNA Library Preparation: M.B. Soares Lab
CDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution i
found through the I.M.A.G.E. Consortium/LLNL
www-bio.llnl.gov/bbrp/image/image.html
Seq primer: M13 Forward.
ACTACTACTACGGTATGGACGTCTGGGGCCAAGGGACCACGGTCACCGTCTTTTCAGCTT
                                               ACCACTACTACGGTTTGGACGTCTGGGGCAAAGGGGACCACGGTCACCGTCTCCTCAGGAT
                                                                                                                                                          CTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGCC---TGTTGGGTGACTATG
                                                                                                                                                                                                                TCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATCTGCAAATGAACAGCCTGAGAA
                                                                                                                                                                                                                                                  TCACCATCTCCAGAGACACTTCCAAGAACACGGTGTATCTAAAAATGAACAGCCTGAGAA
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/ Librus - Lyppe - Lymph |
/ Cell _ Lype - Lymph |
/ Cell _ Line - WACC85 |
/ Lab _ Tost = "PH10B (LTI) |
/ Clone _ Lib = "NH MGC 36"
/ Clone _ Lib = "NH MGC 36"
/ Constructed from size fractionated cytoplasmic mRNA (0.5-1.5kb) . Directionally cloned . Cells provided by M. Staudt, Ph.D. Library preparation by Maria de Fati Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
                                                                                                        TGTATTACTGTGCGAAAGCCCCTGGTCCTACGGTGACTTCACCTT
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Pred. No. 1.3e-70;
0; Mismatches 57
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Homo

sapiens linear

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 536)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
BQ709375
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Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer: M13 Forward.
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/(Cell_line="IDH10B (LTI)"
/(Lone_lib="NIH_MGC_36"
/(clone_lib="NIH_MGC_36"
/(note="Vector: pT7T3-Pac; Site_1: Not1; Site_2: Eco RI;
/(note="Vector: pT7T3-Pac; Site_1: Not1; Site_2: Not1; Site_2: Not1; Site_2: Not1; Site_2: Not1; Site_2: Not1; Site_2: Not1; Si
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/cell_type="germinal center B
/cell_line="MGC85"
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/db_xref="taxon:9606"
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|mol_type="mRNA"
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Pred. No. 3.2e-70;
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Tissue Procurement: Dr. Mark Watson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 881)
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BQ709375
BQ709375.1 GI:21848274
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Homo sapiens
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/lab host="DH108 (phage-resistant)"
/clone_lib="NIH MGC 113"
/clone_lib="NIH MGC 113"
/note="Organ: spleen; Vector: pOTB7; Site_1: XhoI; Site_2: /note= /
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Tissue Procurement: Dr. Mark Watson
CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM2385 row: g column: 16
High quality sequence stop: 401.
Location/Qualifiers
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BQ709399
BQ709399.1 GI:21848298
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1 (bases 1 to 979)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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                                    CCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCTCAG
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61 GAAATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC 120	DЪ
70 GATATIGIGITGACGCAGICTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC 129	Ş
Query Match 33.7%; Score 309; DB 10; Length 424; Best Local Similarity 93.4%; Pred. No. 3.9e-69; Matches 337; Conservative 0; Mismatches 15; Indels 9; Gaps 1;	Que Bes Mat
//Objection noise september // (Including type = mRNA)   // (ab xref="taxon:9606"	BASE COUNT
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Tel; +55-11-2704922  Fax: +55-11-2707001:  Email: asimpson@ludwig.org.br  Email: asimpson@ludwig.org.br  This sequence was derived from the FAPESP/LICR Human Cancer Genome  Project. This entry can be seen in the following URL  (http://www.ludwig.org.br/scripts/gethtml2.pl?t1=II3&t2=IL3-ET0115-  071100-302-D08&t3=2000-11-07&t4=1)	
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_	VERSION KEYWORDS SOURCE ORGANISH
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371 ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC 430	đđ
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                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov

Eco RI site shown at the beginning of the sequence.

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

cDNA Library Preparation: M.B. Soares Lab

cDNA Library Arrayed by: M.B. Soares Lab

DNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution information

found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AW405972
UI-HF-BLO-acv-a-01-0-UI.r1 NIH NIMAGE:3060360 5', mRNA sequence
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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   Conservative
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(1.5-2.5kb). Directionally cloned. Cells provided by Louis
M. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
29 a 169 c 145 g 123 t
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.B. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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603042678F1 NIH_MGC_116 Homo
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol type="mRNA"
/db_xref="teaxon:9606"
/clone="TMAGE:5183010"
/lab_host="DH108"
/clone lib="NIH MGC 116"
/rote="Organ: pooled colon, kidney, stomach; Vector:
/note="Organ: pooled colon, kidney (destroyed); RNA
pCMV-SPORT6; Site_1: Not1; Site_2: EcoRV (destroyed); RNA
source anonymous pool of 3 colons, age 26 yo male, 49 yo
female, 71 yo male colon; 46 yo male kidney, and pool of 2
stomachs, 62 yo male and 70 yo female. Library is
oligo-dT primed and directionally cloned (EcoRV site is
olsetroyed upon cloning). Average insert size 1.4 kb,
insert size range 1-3 kb. Library is normalized and
enriched for full-length clones and was constructed by C.
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                                                                                                                  Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, 1
201203, P. R. China
Tel: 86-21-50801919 (ex.45)
Fax: 86-21-50801922
                                                                                                                                                                                                                                            Xu,X., Huang,J., Xu,Z., Qian,B., Zhu,Z., Yan,Q., Cai,T., Zhang,X., Xiao,H., Qu,J., Liu,F., Huang,Q., Cheng,Z., Li,N., Du,J., Hu,W., Shen,K., Lu,G., Pu,G., Zhong,M., Xu,S., Gu,W., Huang,W., Zhao,X., Hu,G., Gu,J., Chen,Z. and Han,Z.
Insight into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
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1 (bases 1 to 433)
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AV647117 GLC Homo sapiens
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                                                                         l: hanzg@chgc.sh.cn
clone is available at Cr
Location/Qualifiers
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             /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                           Contact: Zeguang Han
Chinese National Human Genor
351 Guo Shoujing Road, Zhang
201203, p. R. China
Tel: 86-21-50801919 (ex.45)
Fax: 86-21-50801922
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                                                                                                                                       Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 440)

Xu,X., Huang,J., Xu,Z., Qian,B., Zhu,Z., Yan,Q., Cai,T., Zhang,X., Xiao,H., Qu,J., Liu,F., Huang,Q., Cheng,Z., Li,N., Du,J., Hu,W., Xiao,H., X., Lu,G., Fu,G., Zhong,M., Xu,S., Gu,W., Huang,W., Zhao,X., Hu,G., Gu,J., Chen,Z. and Han,Z.

Insight into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
                                                                                                                     21625106
11752456
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 Email: hanzg@chgc.sh.cn
This clone is available
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/note="Vector: pBluescript sk(-);
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/tissue_type="corresponding
/dev_stage="Adult"
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Zhangjiang Hi-Tech Park,
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       Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the sequence.
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information
                                                                                                                                                                                                                                       AW405787 504 bp mRNA UI-HF-BLO-abp-e-02-0-UI.rl NIH_MGC_37 Homo IMAGE:3057482 5', mRNA Bequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 504)
                                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                                                  Homo sapiens
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/dev_stage="Adult"
/lab_host="SOLR"
/clone_lib="GLC"
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/clone="GLCAUG10"
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|mol_type="mRNA"
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Unpublished
Contact: Robert Strausberg,
Email: cgapbs-r@mail.nih.go
Tissue Procurement: Louis M
                                            Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 734)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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'db_xref="taxon:9606"
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                                                                                                                                             GI:13918145
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Pred. No. 8.6e-69;
D; Mismatches 12
M. Staudt,
                                                                                                                                                                                           734 bp
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                                                                                                                                                                                sapiens
                                                                                                                                                                                            mRNA
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M.D.,
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365 ---ACGTTCGGCCAAGGGACCAAGGTGGAAATCAAACGAACTGTGGGTGCACCATCTGTC
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/cell_type="germinal center B cells"
/cell_line="McC85"
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/lab_host="DH10B (LTI)"
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/note="Vector: pT7T3-Pac; Site_1: Not1; Site_2: Bco RI;
/constructed from size fractionated cytoplasmic mRNA
(1.5-2.5kb). Directionally cloned. Cells provided by Louis
M. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
17 a 147 c 127 g 113 t
                                                                        Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                            cDNA clone IMAGE: 4763021 5',
Gene Collection
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DEFINITION
ACCESSION
VERSION
                                                                 SOURCE
ORGANISM
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AV646708
                                                                                                                          KEYWORDS
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Best Local Similarity
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AV646708.1
EST.
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     Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                           Homo sapiens (human)
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Location/Qualifiers
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Plate: LLCM1618 row: a column: 06
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                                                                                                                                                                                                                                                                                                                                                                   T 415
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="primary B-cells from tonsils (cell line)"
/lab host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/note="forgan; B-cells; Vector: pOTB7; Site_1: XhoI;
/note="forgan; B-cells; Vector: potB7; Site_1: XhoI
                                                                                                                                                                                              GLC Homo sapiens
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 1e-68;
0; Mismatches 12;
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        Euteleostomi; Homo.
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Ś 밁 Ś 밁 Ş 밁 8 문 Ś 밁 Ş 밁

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BASE COUNT
ORIGIN
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Best Local S
Matches 337
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MEDLINE
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AGENCOURT_7975592 NIH_MGC_113 Homo 5', mRNA sequence.
EQ706634
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Insight into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: hanzg@chgc.sh.cn
This clone is available at CHGC in
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Zeguang Han
Chinese National Human Gen
351 Guo Shoujing Road, Zha
201203, P. R. China
Tel: 86-21-50801919(ex.45)
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Similarity 93.4%;
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="GLCAQE01"
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/note="Vector: pBluescript sk(-); Site_1:
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1 (bases 1 to 926)
NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCR0288 row: h column: 07
High quality sequence stop: 536.
Location/Qualifiers
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Tissue Procurement: Dr. Mark Watson
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Mark Watson
CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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/mol type="mRNA"
/mol type="mRNA"
/db xref="taxon:9606"
/clone="IMAGE:6214925"
/lab host="PH10B (phage-resistant)"
/clone_lib="NIH MGC_113"
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_113 Homo sapiens cDNA clone IMAGE:6214925
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Qy 430 T 430

Db 414 T 414

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Search completed: September 17, 2003, 04:21:49
Job time: 1938 secs

Title: Perfect score:

US-08-657-449-13
918
1 GAATTCATGAAAAAACCGC.....ATCACCATTAGTGAAAGCTT 918

September 17, 2003, 03:02:49; Search time 2893 Seconds (without alignments)
10523.453 Million cell updates/sec

Run on:

OM nucleic - nucleic search, using sw model

GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd

Sequence:

Scoring table:

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. / Cgm2_6/ptodata/2/pna/US103B_COMB.seq:
. / Cgm2_6/ptodata/2/pna/US1001C_COMB.seq:
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/cgn2_6/ptodata/2/nna/ns100A_COMB.seq:*
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

Pending\_Patents\_NA

6/prodata/2/pna/PCTUS\_COMB.seq:\*
6/prodata/2/pna/PCTUS\_COMB.seq.old:\*
6/prodata/2/pna/USO6\_COMB.seq:\*

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Total number of hits satisfying chosen parameters:

33363688 seqs, 16581889874 residues

Searched:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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US-09-782-397-13	US-09-697-561A-13	US-09-697-561-13	US-08-657-449-13		STIMMARTES
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RESULT 1
US-08-657-449-13
                                                                                                                                         Sequence 13, Application US/08657449

GENERAL INFORMATION:

APPLICANT: Dan, Michael D.

APPLICANT: Maiti, Pradip K.

APPLICANT: Kaplan, Howard A.

APPLICANT: Kaplan, Howard A.

TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLECTIDES ENCODING TITLE OF INVENTION: FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND TITLE OF INVENTION: DETECTION OF CANCERS

NUMBER OF SEQUENCES: 18
ZIP: 94304-1018

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/657,449
FILING DATE:
                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: MOTTISON &
STREET: 755 Page Mill
CITY: Palo Alto
STATE: CA
COUNTRY: USA
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US-09-782-397A-16
US-09-782-397B-16
US-09-782-397B-16
US-09-782-397C-16
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TELEX: 706.141
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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Best Local Similarity 100.
Matches 918; Conservative
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ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 11FORMATION:
TELECOMMUNICATION 11FORMATION:
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RESULT 2
US-09-697-561-13
; Sequence 13, Application US/09697561
; GENERAL INFORMATION:
APPLICANT: Dan, Michael D.
Maiti, Pradip K.
Howard A.
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Best Local S
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TELEX: 706141
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleis: single
                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/697,561
FILING DATE: 26-Oct-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
                                                                Local Similarity
nes 918; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Fo
STREET: 755 Page Mill R
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Lebnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dan, Michael ...
Maiti, Pradip K.
Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
                                                                                                                                         NAME/KEY: CDS
LOCATION: join(1..906, 913..918)
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
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                STATE: CA
COUNTRY: U
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                                                                           Score 918; DB 30;
Pred. No. 2.5e-225;
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        RESULT 3

US-09-697-561A-13

; Sequence 13, Application US/09697561A

; GENERAL INFORMATION:

; APPLICANT: Viventia Biotech Inc.

; APPLICANT: Dan, Michael D

; APPLICANT: Maiti, Pradip K

; APPLICANT: Kaplan, Howard A.

; TITLE OF INVENTION: Antigen Binding Fr;

; TITLE OF INVENTION: Nucleotides Encod:
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; LOCATION: (913)..(918)
; OTHER INFORMATION:
US-09-697-561A-13
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CURRENT APPLICATION NUMBER: U5/09/697,561A
CURRENT FILING DATE: 2000-10-26
PRIOR APPLICATION NUMBER: 08/862,124
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin version 3.1
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 918; Conser
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ORGANISM: homo sapien
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NAME/KEY: CDS
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                                                SEQUENCE DESCRIPTION: SEUS-09-782-397-13
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                Query Match
Best Local (
   Matches
                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: LIBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PACENTIN Release #1.0, Version #1.

CURRENT APPLICATION DATA:

APPLICATION UNMBER: US/09/782,397

FILING DATE: 13-Feb-2001

CLASSIFICATION: CIDKNOWN>

PRIOR APPLICATION UNMBER: 08/862,124

FILING DATE: 1997-05-22

ATTORNEY/AGENT INFORMATION:

NAME: Lehnhardt, Susan K.

REGISTRATION UNMBER: 33,943

REFERENCES DOCKET NUMBER: 31608-20001.20

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS: & Foer:
ADDRESSEE: MORTISON & Foer:
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Maiti, Pradip K.

Kaplan, Howard A.

TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                              FEATURE:
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Palo Alto
                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                           TELEX: 706141
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SEQ ID NO: 13:
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Score 918; DB 55,
Pred. No. 2.5e-225;
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RESULT 5
US-09-782-397A-13
Sequence 13, Application US/09782397A
GENERAL INFORMATION:
APPLICANT: Dan, Michael D.
Maiti, Pradip K.
Kaplan, Howard A.
ITTLE OF INVENTION: APPGIFICALLY
FRAGMENTS, AN
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Maiti, Pradip K.
KADIAN, HOWARD A.
INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
INVENTION: ANTIGEN BINDING FRAGMENTS, NUCLEOTIDES ENCODING
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
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LOCATION: join(1..906, 913..918)
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-09-782-397A-13
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid.
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94304-1018

ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYRE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Vers

SOFTWARE: Patentin Release #1.0, Vers

CURRENT APPLICATION DATA:

APPLICATION WINDER: US/09/782,397A

FILING DATE: 13-Feb-2001

CLASSIFICATION: CUNKNOWN>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEFAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/862,124
FILING DATE: 22-MAY-1997
ATTORNEY/AGENT INFORMATION:
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                            361
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                                               ACACCTCAGATCACTTTCGGCGGAGGGACCCAAGGTGGAGATCAAACGAACTGTGGCTGCA
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RESULT 6
US-09-782-397B-13
; Sequence 13, Application US/09782397B
; GENERAL INFORMATION:
                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Vers

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/782,397B

FILING DATE: 13-Feb-2001

CLASSIFICATION UMBER: 08/862,124

PRIOR APPLICATION UMBER: 08/862,124

PILING DATE: 1997-05-22

APPLICATION INFORMATION:

APPLICATION INFORMATION:

APPLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster LLP
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: CA
     NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
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INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT

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TELEFAX: (650) 494-0792
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AAAAAAAACCGCTATCGCGATCGCAGTTGCACTGGCTGGTTTCGCTACCGTT 60	NAME/KEY: CDS  I LOCATION: (1)(906)  OTHER INFORMATION:  NAME/KEY: CDS  LOCATION: (913)(918)  OTHER INFORMATION:  US-09-782-397C-13  Query Match  Best Local Similarity 100.0%; Score 918; DB 33; Length 918;  Best Local Similarity 100.0%; Pred. No. 2.5e-225;  Matches 918; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	CURRENT APPLICATION NOWBER: US/09/782,397C  CURRENT FILING DATE: 2001-02-13  PRIOR APPLICATION NUMBER: 08/862,124  PRIOR FILING DATE: 1997-05-22  PRIOR FILING DATE: 1997-05-22  PRIOR FILING DATE: 1996-05-22  NUMBER OF SEQ ID NOS: 29  SOPTWARE: PatentIn version 3.1  LENGTH: 918  TYPE: DNA  ORGANISM: homo sapien	RESULT 7 US-09-782-397C-13 Sequence 13, Application US/09782397C GENERAL INFORMATION: APPLICANT: Viventia Biotech Inc. APPLICANT: Dan, Michael D APPLICANT: Maiti, Pradip K APPLICANT: Maiti, Pradip K APPLICANT: Kaplan, Howard A. TITLE OF INVENTION: Antigen Binding Fragments that Specifically Detect Cancer Cells, TITLE OF INVENTION: Nucleotides Encoding the fragments FILE REFERENCE: 830001-2001.2	The Third Continger of the Continue of the Continger of the Continger of the Continger of the Continue of the Cont
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT TITLE OF INVENTION: SPECIFICALLY DEFECT CELLS, NUCLEOTIDES ENCODING THE TITLE OF INVENTION: SPECIFICALLY DEFECT CANCER CELLS, NUCLEOTIDES ENCODING THE INTERIOR OF INVENTION: DETECTION OF CANCERS NUMBER OF SEQUENCES: 18 CORRESPONDENCE ADDRESS: ADDRESSE: MOTIBON & FOOTSET STREET: 755 PAGE MILL ROAD CITY: PAIO ALC STATE: CA COUNTRY: USA ZIP: 94304-10.18 COMPUTER READABLE FORM: MEDIUM TYPE: FLOPBY disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTON DATA: APPLICATION NUMBER: US/08/657,449 FILING DATE: CLASSIFICATION: 424 ATTORNEY/AGENT INFORMATION: NAME: Lehnhardt, Susan K. REGISTRATION UNMBER: 33,943 REFERENCE/DOCKET NUMBER: 31608-20001.00 TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION:	QY 901 CACCATTAGTGAAAGCTT 918	721 721 781 781 841 841	Qy  541 TGTGCAGCCTCTGGATTCCCCTTCAGAAGCTTTGCACTGGGTCCGCCAGGCTCTA 600	Db 361 ACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCA 420  Qy 421 CCATCTGTCTCTGGCGGTGGCGGTTCCGGAGGTGGATCAGGTGGAGGTGGCTCCCAG 480

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TELEX: 706141
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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Best Local
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ACCGTCTCCTCAGGATCCGAACAAAAACTGATCAGCGAAGAAGATCTGAACCATCACCAT
                                                       CTGTTGGGTGACTATGACCACTACGGTTTTGGACGTCTGGGGCAAAGGGACCACGGTC
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Pred. No. 6.5e-225;
0; Mismatches 1;
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RESULT 9
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APPLICANT: Dan, Mi
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECHONE: (650) 813-5600
TELEPHONE: (650) 813-5600
TELEPHONE: (650) 494-0792
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/697,561
FILING DATE: 26-Oct-2000
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESSE: MOORTISON & FOORBER
ADDRESSE: 755 Page Mill Road
CITY: Palo Alto
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MEDIUM TYPE: Floppy disk
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Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID
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                                                                                                                                  GCGCAGGCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAA
                 CAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACT
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TYPE: nucleic acid
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 CAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACT
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APPLICANT: Viventia Biotech Inc.
APPLICANT: Dan, Michael D
APPLICANT: Maiti, Pradip K
APPLICANT: Maiti, Pradip K
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: Antigen Binding Fragments that Specifically Detect Cancernation of Invention Nucleotides Encoding the fragments...
FILE REFERENCE: 300001-2001.1
CURRENT APPLICATION NUMBER: US/09/697,561A
CURRENT APPLICATION NUMBER: 08/862,124
PRIOR APPLICATION NUMBER: 08/862,124
PRIOR FILING DATE: 1997-05-22
NUMBER OF SEQ ID NOS: 29
SOPTWARE: Patentin version 3.1
SEQ ID NO 15
LENGTH: 918
TYPE: DNA
COGANISM: Homo Sapien
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US-09-697-561A-15/c
15-Gequence 15, Application US/09697561A
GENERAL INFORMATION:
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RESULT 11
US-09-182-397-15/c
; Sequence 15, Application US/09782397
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Query Match
Best Local Similarity
Matches 917; Conserv
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Pred. No. 6.5e-225;
0; Mismatches 1;
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TELEX: 706141

INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
US-09-782-397-15
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FILING DATE: 13-Feb-2001
CLASSIFICATION CURROWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: MOTTISON & FOETSTET LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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ACACCTCAGATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCA
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Maiti, Pradip K.
Kaplan, Howard A.
KAPlan, Howard A.
INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING THE
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
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US-09-782-397A-15/c
US-09-782-397A-15/c
; Sequence 15, Application US/09782397A
; GENERAL INFORMATION:
; GENERAL TORON, Michael D.
. APPLICANT: Dan, Michael D.
. APPLICANT: Maiti, Pradip K.
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                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: LIM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/782,397A

FILING DATE: 13-Peb-2001

CLASSIFICATION: cUnknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/862,124

APPLICATION NUMBER: US 08/862,124

APPLICATION NUMBER: US 08/862,124

APPLICATION NUMBER: US 08/862,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster LLP
STREET: 755 Page Mill Road
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kaplan, Howard A.

TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
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                    NAME: Lehnhardt,
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
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               CTGTTGGGTGACTATGACCACTACTACGGTTTGGACGTCTGGGGCAAAGGGACCACGGTC
                                                                                                   GACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGGTGTATCTA
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                                           AAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGC
Conservative
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SEQUENCE DESCRIPTION: SEQ ID NO: 15: US-09-782-397B-15
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US-09-782-397B-15/c
Sequence 15, Application US/09782397B
GENERAL INFORMATION:
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                                                                                                                                                                                                                             Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/782,397B
FILING DATE: 13.-Feb-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,124
FILING DATE: 1997-05-22
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.20
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Maiti, Pradip K.
KADIAN, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS H11, THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES
ENCODING THE
ENCODING THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Dan,
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ACCGTCTCCTCAGGATCCGAACAAAAACTGATCAGCGAAGAAGATCTGAACCATCACCAT
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                                                                                                                                                                                                                                                                                                                                        LENGTH: 918 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                         GCGCAGGCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAA
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STATE: CA
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                                                                                                                                                                                                              Conservative
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Pred. No. 6.5e-225;
0; Mismatches 1;
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RESULT 14
US-09-782-397C-15/c
US-09-782-397C-15/c
Sequence 15, Application US/09782397C
GENERAL INFORMATION:
APPLICANT: Viventia Biotech Inc.
APPLICANT: Dan, Michael D
APPLICANT: Maiti, Pradip K
APPLICANT: Maiti, Pradip K
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: Antigen Binding Fragments that Specifically Detect Cancer Cells,
TITLE OF INVENTION: Nucleotides Encoding the fragments...
FILE REFERENCE: 830001-2001.2
CURRENT APPLICATION NUMBER: US/09/782,397C
CURRENT FILING DATE: 1001-02-13
PRIOR APPLICATION NUMBER: 08/652,124
PRIOR FILING DATE: 1997-05-22
PRIOR FILING DATE: 1996-05-22
NUMBER OF SEQ ID NOS: 29
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US-09-782-397C-15
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Best Local Sim
Matches 917;
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SEQ ID NO 15
LENGTH: 918
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FEATURE:
NAME/KEY:
LOCATION:
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CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,449
FILING DATE:
CLASSIFICATION SUBBER: US/08/657,449
FILING DATE:
NAME: Lehnhardt, Suban K.
REGISTRATION NUMBER: 33,943
REPERENCE/DOCKET NUMBER: 33,943
REPERENCE/DOCKET NUMBER: 31608-20001.00
TELEPHONE: (415) 813-5600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELECOMINICATION UND SEGUENCE: SEGUENCE CHARACTERISTICS:
LENGTH: 867 base pairs
TYPE: nucleic acid
STDANNEDNESS: single
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Best Local Similarity
Matches 863; Conserv
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APPLICANT: Maiti, Pradip K.
APPLICANT: Maiti, Pradip K.
APPLICANT: Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT
TITLE OF INVENTION: SPECIFICALLY DETECT CANCER CELLS, NUCLECTIDES ENCODING THE
TITLE OF INVENTION: FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
TITLE OF INVENTION: DETECTION OF CANCERS
TITLE OF INVENTION: DETECTION OF CANCERS
TORRESPONDENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: MOTILISON & FOCISICE
STREET: 755 Page Mill Road
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                    241 GGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGT
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GGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGT
                                                                                  CAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACT 240
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Search completed: September 17, 2003, 05:12:00 Job time : 2906 secs

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(cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*
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(cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
(cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
(cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
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ALIGNMENTS

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## US-09-104-164-13 SEQUENCE 13, Application US/09194164 GENERAL INFORMATION: Michael D. APPLICANT: Dan, Michael D. APPLICANT: Dan, Michael D. KADIAN, Howard A. KADIAN, Howard A. KADIAN, Howard A. KADIAN, HOWART CAND USE THERROF FOR THE PROPHYLAXIS AND DESCRIPTION: ANTIGEN BINDING FRAGMENTS THAT SPECIFICALLY DEFECT CANCER CELLS, NUCLEOTIDES ENCODING FRACMENTS, AND USE THERROF FOR THE PROPHYLAXIS AND NUMBER OF SEQUENCES: 18 CORPERED OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT SPECIFICALLY DEFECT CANCERS NUMBER OF SEQUENCES: 18 CORPUTES: 18 AD USE THERROF FOR THE PROPHYLAXIS AND NUMBER: 755 Page Mill Road GITY: Palo Alto GITYANDENES: 1 1004 GITANDENES: 1 1004 GITTY: Palo Base pair B GITANDENES: 1 1004 GITANDENES: 1 1004 GITTY: Palo Base pair B GITTY: P

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GENERAL INFORMATION:
APPLICANT: Dan, Michael D.
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Best Local Similarity 99.9%;
Matches 917; Conservative
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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PILING DATE: CURKNOWN>
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.00
TELEPHONE: (415) 813-5600
TELEPHONE: (415) 813-5600
TELEPHONE: (415) 494-0792
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ZIP: 94304-1018

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/99/194,164
PILING DATE: 09-Apr-1999
CLASSIFICATION: CUMPONN
PRIOR APPLICATION DATA:
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Kaplan, Howard A.
TITLE OF INVENTION: ANTIGEN
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison
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TYPE: nucleic acid
STRANDEDNESS: single
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COUNTRY: USA
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                                                                                                            GGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGT
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                                                                      GGCATGCCAGACAGGTTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGT
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SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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Pred. No. 1.5e-257;
0; Mismatches 1;
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RESULT 3
US-09-194-164-16
; Sequence 16, Application US/09194164
; GENERAL INFORMATION:
GENERAL INFORMATION:
COMPUTER READABLE FORM:

WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/194,164
FILING DATE: 09-Apr-199
CLASSIFICATION: cÜnknown>
PRIOR APPLICATION NUMBER: US/08/657,449
FILING DATB: «Unknown>
PRIOR APPLICATION NUMBER: US/08/657,449
FILING DATE: «Unknown>
                                                                                                                                                                                                                                                                                                                  Maiti, Pradip K.
Kaplan, Howard A.

TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
PRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison &
STREET: 755 Page Mill
CITY: Palo Alto
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COUNTRY: USA
ZIP: 94304-1018
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
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LOCATION: join(1..855, 862..867)
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
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REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susar
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   AAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGC
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94.0%;
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Pred. No. 1.8e-223;
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US-09-194-164-18/c
; Sequence 18, Application US/09194164
; GENERAL INFORMATION:
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; SEQUENCE DESCRIPTION: SEQ ID NO: 18: US-09-194-164-18
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                                                                                           Query Match
Best Local Similarity
Matches 862; Conserv
                                                                                                                                                                                                                                                TELEX: 706141
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/194,164
FILING DATE: 09-Apr-1999
CLASSIFICATION: Unknown>
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Kaplan, Howard A.

TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT

SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING

SPECIFICALLY DETECT CANCER THE PROPHYLAXIS AND

FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/657,449
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Lebnhardt, Susan K.
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ADDRESSEE: Morrison & Foerster
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                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 31608-20001.00
TELECOMMUNICATION INFORMATION:
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STATE: CA
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GCGCAGGCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAA 120
                                                 COUNTRY: USA
                                STRANDEDNESS: single
                                                                                                                                                                                                                     LENGTH: 867 base pairs
TYPE: nucleic acid
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TELEFAX: (415) 494-0792
                                                                                               Conservative
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                                                                                                            Score 798; DB 5;
Pred. No. 5.4e-223;
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                                                                                                                                                                               RESULT 5
PCT-US03-14964-4
        APPLICANT: Rother, Russell P
APPLICANT: Geis, David R
TITLE OF INVENTION: T7 Bacteriophage Display of
FILE REFERENCE: 1087-24 PCT
CURRENT APPLICATION NUMBER: PCT/US03/14964
CURRENT FILLING DATE: 2003-07-17
PRIOR APPLICATION NUMBER: US 60/380,318
PRIOR FILING DATE: 2002-05-14
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                Sequence 4, Application PC/TUS0314964; GENERAL INFORMATION:
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Best Local Similarity
Matches 628; Conser
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SEQ ID NO 4
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TYPE: DNA
ORGANISM: human
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                                                                                                            GGTGACTATGACCACTACTACGGTTTGGGACGTCTGGGGCAAAGGGACCACGGTCACCGTC
                                                                                                                                                 AGCAGCCTGAGATCTGAGGACACGGCCATATATTATTGTGCGAGAGGGGATACGATTTTT
                                                                                                                                                                                                                                                          GGGCTTGAGTGGGATGGGATCTTCCCTTTCCGTAATACAGCAAAGTACGCACAACAC
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                                                                                                                                                                                                                                                                                                               GCTTCTGGAGGCACCTTCAACAATTATGCCATCAGCTGGGTGCGACAGGCCCCTGGACAA
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                                                                       TCCTC
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Pred. No. 2.1e-123;
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Best Local S
Matches 361
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ATCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTC
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RESULT

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US-09-194-164-6/c
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/194,164

FILING DATE: 09-Apr'-1999

CLASSIFICATION - UNKNOWN>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/657,449

FILING DATE: -UNKNOWN>
APPLICATION NUMBER: US/08/657,449

FILING DATE: -UNKNOWN>
AFTORNEY/AGENT INFORMATION:

NAME: Lehnhardt, Susan K.

REGISTRATION NUMBER: 33,943

REGISTRATION NUMBER: 33,943

REGISTRATION NUMBER: 31608-20001.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
mes 361; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison &
STREET: 755 Page Mill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                  CITY: Palo Alto
STATE: CA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                           CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACCTTAGCCTGGTACCAGCAGAAA
                          CCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCTCAG
                                                                                                                GACAGGTTCAGTGGCAGTĞGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTGGAG
                                                                                                                                                                                                             CCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCACTGGCATGCCA
                                                                                                                                                                                                                                                                              CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAGAAA
CCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCTCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 450 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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ilarity 100.0%;
Conservative (
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INVENTION: ANTIGEN BINDING FRAGMENTS THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
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| Road
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US-09-194-164-4
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: Dan, Michael D.
                                                                                                                                                                                                       Matches
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INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/657,449
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REGISTRATION NUMBER: 31608-20001.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEPAX: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/194,164
FILING DATE: 09-Apr-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT

SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                       FEATURE:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: Morrison & Foerster
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                                                                                    CTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAGAAA 189
                        CCTGGCCAGGCTCCCAGGCTCCATCTATGGTGCATCCACCAGGGCCACTGGCATGCCA
                                                                                                                                   GATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC
                                                                                                                                                          GATATTGTGTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACC 129
                                                                                                                                                                                                                                                                                                      NAME/KEY:
LOCATION:
CTGGCCAGGCTCCCAGGCTCCATCTATGGTGCATCCACCAGGGCCACTGGCATGCCA
                                                                   CTCTCCTGCAGGGCCAGTCAGAGTGTTAGCAGCAGCTACTTAGCCTGGTACCAGCAGAAA
                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 450 base pairs TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
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Pred. No. 4.7e-95;
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                                                                                                                                                                                                                                 DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Version #1.30
                                                                                                                                                                                                                                 Length 450;
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RESULT 8
US-09-194-164-1
Query Match
Best Local Similarity 89.
Matches 349; Conservative
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                                                                                                                                                                                                                                                                                                   TELEX: 706141
INFORMATION FOR SEQ ID NO: 1:
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APPLICATION NUMBER: US/08/657,449
FILING DATE: <UNATION:
ATTORNEY/AGENT INFORMATION:
NAME: Lehnhardt, Susan K.
REGISTRATION NUMBER: 33,943
REGISTRATION NUMBER: 33,943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/194,164
FILING DATE: 09-Apr-199
CLASSIFICATION: <Unknown>
                                                                                                         LOCATION: 1..543
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 31608-20001.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 543 base pairs
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                TYPE: nucleic acid: STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Morrison & Foerster STREET: 755 Page Mill Road CITY: Palo Alto
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                                                                                                                                                  NAME/KEY: CDS
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ZIP: 94304-1018
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                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 494-0792
                                                                                                                                                                                              linear
  35.3%; Score 324.4; DB 5;
89.3%; Pred. No. 8.4e-85;
tive 0; Mismatches 42;
                                           Length 543;
      Indels
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RESULT 9
US-09-194-164-3/c
; Sequence 3, Application US/09194164
; GENERAL INFORMATION:
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            TELEX: 706141
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 543 base pairs
TYPE: nucleic acid
                                                                                                                                             COMPUTER: IBM PCFCOMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/194,164

FILING DATE: 09-Apr-1999

CLASSIPICATION -CURKNOWN>

PRIOR APPLICATION UMBER: US/08/657,449

APPLICATION UMBER: US/08/657,449

APPLICATION UMBER: 33,943

REFERENCE/DOCKET NUMBER: 33,943

REFERENCE/DOCKET NUMBER: 31608-20001.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Maiti, Pradip K.
KAPlan, Howard A.
TITLE OF INVENTION: ANTIGEN BINDING FRAGMENTS THAT
SPECIFICALLY DETECT CANCER CELLS, NUCLEOTIDES ENCODING
FRAGMENTS, AND USE THEREOF FOR THE PROPHYLAXIS AND
DETECTION OF CANCERS
                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                  TELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
STATE: CA
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                                                                                                  TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TCACCATCTCCAGAGACACTTCCAAGAACACGGTGTATCTAAAAATGAACAGGCTGAGAA
STRANDEDNESS: double
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; SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-194-164-3
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PCT-US03-14964-1
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Best Local S
Matches 375
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Best Local Similarity 89.3%;
Matches 349; Conservative
                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.2 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: T7 Bacteriophage Display FILE REFERENCE: 1087-24 PCT CURENT APPLICATION NUMBER: PCT/US03/14964 CURRENT FILING DATE: 2003-07-17 PRIOR APPLICATION NUMBER: US 60/380,318 PRIOR FILING DATE: 2002-05-14 NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Rother, Russell APPLICANT: Geis, David R
                                                                                                                                                                                                                                                                                                                            TYPE: DN
                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: human
                                                                                                                                                                                                                      Local Sim
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                                     ACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGTAGCAGCTACTTAGCCTGGTACCAGCAG
                                                                                  GCGG---CCGAGCTCACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCC
                                                                                                           GCCGATATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCC
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                ACCCTCTCCTGCAGGGCCAGTCACAGTGTTAGCAGGGCCTACTTAGCCTGGTACCAGCAG
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                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                    34.7%;
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PCT-US03-26232-145
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TITLE OF INVENTION: CHEMO-ATTRACTANT PROTEIN-1 (MCP-1) AND USES THEREOF
TITLE OF INVENTION: CHEMO-ATTRACTANT PROTEIN-1 (MCP-1) AND USES THEREOF
FILE REFERENCE: ABGENIX.091VPC
CURRENT FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: US 60/404,802
PRIOR APPLICATION NUMBER: US 60/404,802
PRIOR FILING DATE: 2002-08-19
NUMBER OF SEQ ID NOS: 149
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 145
SEQ ID NO 145
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 145, Application PC/TUS0326232
GENERAL INFORMATION:
APPLICANT: ABGENIX, INC.
APPLICANT: GUDAS, Jean M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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GACTATGACCACTACGGTTTGGACGTCTGGGGCAAAGGGACCACGGTCACCGTCTCC
                                                                       AGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGATCAGAGCCTGTTGGGT
                                                                                                                                          AAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGGTGTATCTAAAAATGAAC 729
                                                                                                                                                                                                                                                                                                                           GAGCAGTCGGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCTCCTGTGCAGCG
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                                                                                                                       AAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGCTGTATCTGCAAATGAAC 240
                                                                                                                                                                                      CTGGAGTGGGTGGCAGTTATATGGTATGATGGAAATAATAATAATACTATGCAGACTCCGTG 180
                                                                                                                                                                                                                                                          TCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTCCAGGCAAGGGG
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                                                    AGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGATAGCAGCTCGT-----
                                                                                                                                                                                                                    CTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCACTAAATACTACGCAGACTCCGTG 669
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LIANG, Meina L.
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Pred. No. 5.5e-73;
0; Mismatches 32;
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Sequence 42, Application PC/TUS0321304

GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO ANTI-PLATELET TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS THEREOF FILE REFERENCE: 053893-5050WO
CURRENT FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: ECT/US03/21304
CURRENT FILING DATE: 2003-07-03
PRIOR REPLICATION NUMBER: 60/394,356
PRIOR REPLICATION NUMBER: 60/394,356
PRIOR FILING DATE: 2003-07-03
PRIOR FILING DATE: 2003-09-18
NUMBER OF SEQ ID NOS: 153
SOPTWARE: Patentin version 3.2
SEQ ID NO 42
                                                                                                                                                                                                  RESULT 13
US-09-610-551B-2
; Sequence 2, Application US/09610551B;
; GENERAL INFORMATION:
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PCT-US03-21304-42
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Best Local Similarity
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TYPE: DNA
ORGANISM: Homo sapiens
NUMBER OF SEQUENCES: 70

CORRESPONDENCE ADDRESS:
ADDRESSE: The Scripps Research Institute
STREET: 10666 North Torrey Pines Road, TPC8
CITY: La Jolla
STATE: CA
                                                                                                           APPLICANT: Barbas, Carlos F, III
Burton, Dennis R
Lerner, Richard A
TITLE OF INVENTION: METHODS FOR PRODUCING ANTIBODY LIBRARIES
USING UNIVERSAL OR RANDOMIZED IMMUNOGLOB
                                                                                                                                                                                                                                                                                                                        290
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                                                                                                                                                                                                                                                                                                                                                                                                 CCTGAAGATTTTGCAGTGTATTACTGTCAGCAGTATGGTGGCTCACCTC-
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Pred. No. 9.6e-73;
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RESULT 14
PCT-US21304-4
; Sequence 4, Application PC/TUS0321304
; GENERAL INFORMATION:
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Best Local Similarity
Matches 317; Conserv
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 646 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE DOCKET NUMBER: TS:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 858-554-2937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER: US 07/826,623
FILING DATE: 27-JAN-1992
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                                                                                                                                                TTCGGCGGAGGACCAAGGTGGAGATCAAACGAACTGTGGCTGCACCATCTGTCT 430
                                                                                                                                                                                    GATTTTGCAGTGTACTACTGTCAGCAGTATGGTGGCTCACCGTGG------
                                                                                                                                                                                                                       GATTTTGCAGTGTATTACTGTCAGCAGTATGGTAGCTCACCTCAGACACCTCAGATCACT 375
                                                                                                                                                                                                                                                             TCCAGTGGCAGTGGGTCTGGGACAGACTTCACTCTCACCATCAGCAGACTGGAGCCTGAA
                                                                                                                                                                                                                                                                                                TTCAGTGGCAGTGGGTCCGGGACAGACTTCACTCTCACCATCAGTAGACTGGAGCCTGAA 315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/174,674
FILING DATE: 28-DEC-1993
APPLICATION NUMBER: US 08/300,386
FILING DATE: 02-SEP-1994
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ilarity 89.3%;
Conservative
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Pred. No. 2.7e-71;
D; Mismatches 23
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APPLICANT: Siegel, Donald L.

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TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS REI
TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS THE
FILE REFERENCE: 053893-5050MO
CURRENT APPLICATION NUMBER: PCT/US03/21304
CURRENT FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 60/394,356
PRIOR FILING DATE: 2003-07-03
PRIOR PILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 60/411,694
PRIOR FILING DATE: 2003-09-18
NUMBER OF SEQ ID NOS: 153
SOFTWARE: Patentin version 3.2
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LENGTH: 351
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                         Query Match
Best Local Similarity
Matches 283; Conserv
                                                                                                                                                                                                                                                                 SEQ ID NO 5
LENGTH: 351
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Best Local Similarity 95.6%;
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CURRENT FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 60/394,356
PRIOR FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 60/411,694
PRIOR APPLICATION NUMBER: 60/411,694
PRIOR FILING DATE: 2003-09-18
RUMBER OF SEQ ID NOS: 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS THEREOF FILE REFERENCE: 053893-5050WO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Siegel, Donald L.
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ORGANISM: Homo sapiens
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Search completed: September 17, 2003, 05:16:57 Job time : 283 secs	241 CTGCAAATGAACAGCCTGAGACCCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGT 297	718 CTAAAATGAACAGCCTGAGAACTGAGGACACGGCTGTCTATTACTGTGCGAGAGAT 774	181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240	658 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACACTTCCAAGAACACGTGTAT 717	121 CCAGGCAAGGGGCTGGAGTGGCAGTTATATCATATGATGAAGTAATAAATA	598 CTAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGGATGG	61 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAACTATGCTATGCACTGGGTCCGCCAGGCT 120